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121
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Pred. No. is the number of results predicted by chance б have

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	AX063467	RESULT 1
and dumas milne Edwards. J.B. Apolipoprotein a-iv-related protein: polypeptide, polynucleotide	Yen.F. Denison.B. Bour.B. Bibain B. Bourmeleret I Duclert A	Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens	human.		AX063467.1 GI:12541237	AX063467	•	AX063467 5381 bp DNA linear PAT 24-JAN-2001		

ALIGNMENTS

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Patent: wO 0100803-A 4 04-JAN-2001;
GENSET (FR)
                                                                                         5381 bp
Sequence 4 from Patent WO02053734.
AX469879
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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3214. .3232
/note="17-41-250.mis complement"
3941. .5381
/note="3'regulatory region"
a 1434 c 1709 g 1132 t 2 others
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/note="17-41-250.mis"
3201. 3225
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/db_xref="taxon:9606"
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1613. .1724
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/note="17-42-319.mis complement"
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/note="17-41-250.probe"
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Query Match 99.7%; Score 120.6; DB 6; Best Local Similarity 100.0%; Pred. No. 2.6e-27; Matches 121; Conservative 0; Mismatches 0;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 81001)
                                                                                                                                 AX063464.1 GI:12541234
                                                                                                                                                     Sequence 1 from Patent W00100803 Ax063464
                                                              Homo sapiens
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Apolipoprotein a-iv-related protein: polypeptide, polynucleotide sequences and biallelic markers thereof patent: WO 0100803-A 1 04-JAN-2001;
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/note-"20-828.rp complement"
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45328. .45347
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/note="20-828-311.mis complement"
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/note-"20-828-311.probe"
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/db_xref-"taxon:9606"
/note="20-842.pu"
45423. .45441
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Sequence 1 from Patent WO02053734.
AX469876
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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45863. .45883
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45430. .45454
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/db_xref="taxon:9606"
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12947. .129
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77166. .77185
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            121 C 121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P., Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y. Direct Submission
Submitted (17-MAR-2000) Masahira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); Kitasato Univ., I-15-1 Kitasato, Sagamihara, Kanagawa 228-8555, Japan (E-mail:hattori@gsc.riken.go.jp, URL:http://hgp.gsc.riken.go.jp/, Tel:81-42-778-9923, Fax:81-42-778-9924)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens chromosome 11 clone CTC-227C10 map 11q, WORKING DRAFT SEQUENCE, 18 unordered pieces.

AP001480
NOTE: This is a 'working draft' sequence. It currently consists of 18 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           On May 30, 2000 this sequence version replaced qi:7288164.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 115027)
Hattu.i,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P.,Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y.
Homo sapiens 115,027 genomic DNA of 11q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Published Only in DataBase (2000)
2 (bases 1 to 115027)
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                                                                                                                                                                 Center clone name: CTC-227C10

Center clone name: CTC-227C10

Sequencing vector: PCR products; 100% of reads Chemistry: Dye-terminator ET-amersham; 100% of reads Assembly program: Phrap; version 0.990329

Consensus quality: 107728 bases at least Q40

Consensus quality: 110501 bases at least Q30

Consensus quality: 111810 bases at least Q30

Consensus quality: 111810 bases at least Q30

Consensus quality: 111810 bases at least Q30

Consensus quality: 10720 bases; sum-of-contigs

Quality coverage: 6.70x in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                web site: http://hgp.gsc.riken.go.jp/
Contact: hattori@gsc.riken.go.jp
Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Center: RIKEN Genomic Sciences Center(GSC)
Center code: RIKEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center project name: HumDraftll
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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as soon as it is available and the accession number will be

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FEATURES
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30376 40528 contig of 10434 bp in length
40629 50205 contig of 9577 bp in length
50306 59617 contig of 9577 bp in length
50306 59617 contig of 9912 bp in length
59718 66750 contig of 9912 bp in length
66851 73847 contig of 6997 bp in length
79745 85653 contig of 5697 bp in length
79745 91241 contig of 5488 bp in length
91342 96449 contig of 5078 bp in length
91342 96419 contig of 5179 bp in length
91342 96419 contig of 2179 bp in length
100745 102923 contig of 2179 bp in length
103024 105950 contig of 2277 bp in length
103024 105950 contig of 2339 bp in length
103034 105950 contig of 2539 bp in length
103039 11577 contig of 2539 bp in length
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106051 108938: contig of 2888 bp
108939 109038: gap of 100 bp
109039 111577: contig of 2539 bp
111578 111677: gap of 100 bp
111578 111487: contig of 1810 bp
113488 113487: can of
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100 bp

100644: contig of 4125 b

100645 100744: gap of 100 bp

100745 102923: contid of 102923
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102924 103023: gap of 100 bp
103024 105950: contig of 2927 bp
105951 106650: ~~~ ~~
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79745 85653: contig of 5909 bp in length

85654 85753: gap of 100 bp

85754 91241: contig of 5488 bp in length

91242 91341: gap of 100 bp

91342 96419: contig of 5078 bp in length

96420 96519: gap of 100 bp

96420 96519: gap of 100 bp
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66851 73847: contig of 6997 k
73848 73947: gap of 100 bp
73948 79644: contig of 5697 k
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59618 59717: gap of 100 bp
59718 66750: contig of 7033 bp
66751 66850: gap of
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15682 15781: gap of 100 bp
15782 30275: contig of 14494 bp in length
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                                    /note="assembly_fragment"
15782. .30275
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                                                                                                                                                                                                               /chromosome="11"
                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
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50205: contig of 9577 k
305: gap of 100 bp
59617: contig of 9312 k
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40528: contig of 10153 bp in length
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TITLE
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         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 175582)

Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P., Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y. Homo sapiens 175,582 genomic DNA of 11q

Published Only in DataBase (2000)

2 (bases 1 to 175582)

Hattori,M., Ishii,K., Toyoda,A., Taylor,T.D., Hong-Seog,P., Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y.
                                                                                                                                                                                                                                                                         Homo sapiens chromosome 11 clone CTC-270C21 map 11q, WORKING DRAFT SEQUENCE, 29 unordered pieces.

AP001481
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HTG; HTGS_PHASE1; HTGS_DRAFT.
Homo sapiens DNA, clone:CTC-270C21.
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59718. .66750
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106051. .108938
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96520. .100644
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/note-"assembly_fragment"
79745. .85653
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109039, .111577
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On May 30, 2000 this sequence version replaced gi:7288175.
                                                                          Sequence updated (26-May-2000)

* NOTE: This is a 'working draft' sequence. It currently

* consists of 29 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence.
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                                             as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Web site: http://hgp.gsc.riken.go.jp/
Contact: hattori@gsc.riken.go.jp
------Project Information
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42440 42539: gap of 100 bp
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53422 53521: gap of 100 bp
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16728. .30620
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           Shaikh,T.H., Budarf,M.L., Celle,L., Zackai,E.H. and Emanuel,B.S. Clustered 11q23 and 22q11 breakpoins ad 3:1 meiotic malsegregations in multiple unrelated t(11;22) families
Am. J. Hum. Genet. (1999) In press
2 (bases 1 to 188971)
14u,P. and Roe,B.A.
                                                                                                                                                                                                                                                             Homo sapiens chromosome 11 clone 442e11 from RPCI11 Library map 11q23. complete sequence.
Direct Submission
                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 188971)
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85624. .94708
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165355. .167624
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104177. .112629
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126797. .132422
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160762. .163242
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151526. .155432
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148651. .151425
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138021. .142190
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3 (bases 1 to 188971)
3 (bases 1 to 188971)
Hu,P. and Roe,B.A.
Direct Submission
Submitted (25-AUG-1999) Department Of Chemistry And Biochemistry,
Submitted (25-AUG-1999) Department Of Chemistry And Biochemistry,
Submitted (25-AUG-1999) Department Of Chemistry And Biochemistry,
2 (bases 1 to 42479)
Yang, L., Hu, P., Wu, H., Tapia-Paez, I. and Roe, B.A.
Direct Submission
Submitted (18-JUL-2000) Department Of Chemistry And Biochemistry,
The University Of Oklahoma, 620 Parrington Oval, Room 208, Norman
                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens chromosome 22q11 cl
AC074203
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5 (bases 1 to 188971)
Hu,P. and Roe,B.A.
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                                                                                                            This clone is a chimeric cosmid from a patient with t(11;22) translocation and it is from the derivative chromosome 11. T are sequences from both chromosome 11 and 22 in this cosmid
                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 42479)
Yang,L., Hu,P., Wu,H., Tapla-Paez,I. and Roe,B.A.
Homo sapiens Chromosome 22q11 Cosmid cos4
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On Aug 25, 19
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/Clone-"442e11 from RPCI11 Library"
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/db_xref="taxon:9606"
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Direct Submission
Submitted (13-OCT-2000) Department Of Chemistry And Biochemistry,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yang,L., Hu,P., Wu,H., Tapia-Paez,I. and Roe,B.A. Direct Submission
Center Code: PGABERK.

* NOTE: This is a 'working draft' sequence. It currently

* Consists of 4 contigs. The true order of the pieces

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence

* as soon as it is available and the accession number will
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OK 73019, USA
3 (bases 1 to 42479)
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                                                                                                                                                                                                        Draft
                                                                                                                                                                                                                                                                    Submitted (27-FEB-2002) Berkeley PGA, Lawrence Berkeley National
                                                                                                                                                                                                                                                                                        Direct Submission
                                                                                                                                                                                                                                                                                                           Cheng, J. -F.F.
                                                                                                                                                                                                                                                                                                                                                                           Unpublished
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                                                                                                                                                                                                                                                                                                                                    Martin,J., Hosseini,R., Peng,Y., Peng,Z., Rubin,E.M. and
                                                                                                                                                                                                                                                                                                                                                                                                                       Martin, J , Hosseini, R., Peng, Y., Peng, Z.,
                                                                                                                                                                                                                                               Laboratory, 1 Cyclotron rd., Berkeley, CA 94720, United States of
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Aug 1, 2000 this sequence version replaced gi:9441829
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/clone="cos4"
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/db_xref="taxon:9606"
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Pred. No. 8.1e-27;
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AC118574
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AC118574
Assembly program: Phrap version 0.990329.

* NOTE: This is a 'working draft' sequence. It currently consists of 10 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.
                                                                                                                                         Sequencing vector: Plasmid; pUC18 Chemistry: Dye-terminator Big Dye
                                                                                                                                                                                                            Comparative analysis and ordering by homology are available here: http://pga.lbl.gov/cgi-bin/search_cvcgd?type-n&value-APOAl Funding agent: Programs for Genomic Applications (NHLBI) Contact: 'Jody Schwartz' jrschwartz@lbl.gov
                                                                                                                                                                                                                                                                                                                                              Submitted (19-APR-2002) Genome Sciences, Lawrence Berkeley National Laboratory, I Cyclotron Rd., Berkeley, CA 94720, USA Draft Sequence Produced by Berkeley PGA Web site: http://pga.lbl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Martin, J., Hosseini, R., Peng, Y., Peng, Z., Rubin, E.M. and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cheng, J. -+
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                                                                                                                                                                                           Summary Statistics
                                                                                                                                                                                                                                                                                                      Additional information:
                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
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Mammalia; Eutheria; Primates; Strepsirhini; Lemuridae; Lemur.
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/clone="RP43-68H20"
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48002: gap of unknown length
181412: contig of 133410 bp in length
                                                                                                                                                                                                                                                                                                                              PGABERK
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97.5%;
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Pred. No. 2.
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Db 71391 TTTTGGCTTGCCTCCTCAGAGGCCCTACGCATGGAGTTCAGCCTTTCTCCATGGGGAGA 71450
                                                                                                                                                  BASE COUNT
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AX323766/c
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Local Similarity 79.7%;
                                                                                 Local Similarity
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                                                                    63;
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                                                                                                                                                                                                                                                                                                                                          synthetic construct.
synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 254 from Patent W00192565. AX323766
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                                                                                                                                                                                                                                                                                                                         artificial sequences.
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                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
/note="chemically treated genomic DNA (Homo sapiens)"
176 c 1481 g 2603 t
                                                                                                                                                                                                            Location/Qualifiers
1. .5377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Lemur catta"
/db_xref="taxon:9447"
/clone="LB2-277C3"
41726 c 41047 g 52821 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1. .191656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             102420: contig of 27874 bp in length 102520: gap of unknown length 191656: contig of 89136 bp in length
                                                                                 44.18;
77.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       contig of 3277 l
gap of unknown
contig of 2508 l
gap of unknown
contig of 3209 l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gap of unknown length
contig of 29252 bp in length
gap of unknown length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gap of unknown length
contig of 10569 bp in length
gap of unknown length
contig of 14878 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gap of unknown contig of 4044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gap of unknown contig of 6009
                                                                                 Score 53.4; DB 6; Length 5377; Pred. No. 3.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 80.8; DB 2; Pred. No. 8.1e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bp in length bp in length
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length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   901 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 191656;
                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                          PAT 07-JAN-2002
                                                                0;
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                                                                Gaps
                                                                0;
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В δÃ

SOURCE

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9 GCTTGCTCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCCAAATCTY 68

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JOURNAL REFERENCE
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ORGANISM
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AC118580
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Best Local Similarity Matches 67; Conserv
                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUTHORS
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                                                                                                                                                                                                                           source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   101 CTGGCCCAGATGCCTCTTGCC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assembly program: Phrap version 0.990329.

* NOTE: This is a 'working draft' sequence. It currently consists of 6 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Comparative analysis and ordering by homology are available here: http://pga.lbl.gov/cgi-bin/search_cvcgd?type=n6value=APOAl Funding agent: Programs for Genomic Applications (NHLBI) Contact: 'Jody Schwartz' jrschwartz@lbl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory, 1 Cyclotron Rd., Berkeley, (
Draft Sequence Produced by Berkeley PGA
Web site: http://pga.lbl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (19-APR-2002) Genome Sciences, Lawrence Berkeley National Laboratory, 1 Cyclotron Rd., Berkeley, CA 94720, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oryctolagus cuniculus.
Oryctolagus cuniculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AC118580 233619 bp DNA linear HTG 19-APR Oryctolagus cuniculus clone LB1-82E16, WORKING DRAFT SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing vector: Plasmid; pUC18 Chemistry: Dye-terminator Big Dye
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Martin, J., Hosseini, R., Peng, Y., Peng, Z., Rubin, E.M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HTG; HTGS_PHASE1; HTGS_DRAFT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Summary Statistics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Additional information:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center Code: PGABERK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Martin,J., Hosseini,R., Peng,Y., Peng,Z., Rubin,E.M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cheng, J.-F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      be preserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                as soon as it is available and the accession number will
                                                                                                                         60029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (bases 1 to 233619)
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                                                                                                                                                                                                                                                                                                                                                                   49403
49503
127551
127651
127651
208759
208859
                                                                                                                                                                                                                                                                                              211995
223626
                                                                                                                                                                                                                                                                                                                                               211895
  Conservative
                                                                                                                      /organism="Oryctolagus cuniculus"
/db_xref="taxon:9986"
/clone="Lb1-82E16"
/ 57107 c 56463 g 59513 t 50
                                                                                                                                                                                                                   1. .233619
                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                      49402: contig of 49402 bp in length
49502: gap of unknown length
127550: contig of 78048 bp in length
127650: gap of unknown length
208758: contig of 81108 bp in length
208758: contig of 81108 bp in length
208858: gap of unknown length
211894: contig of 3036 bp in length
211994: gap of unknown length
223625: contig of 11631 bp in length
223775: gap of unknown length
                       38.5%;
                                                                                                                                                                                                                                                                       233619: contig of 9894 bp in length

    Mismatches

                    Score 46.6; DB 2; Length 233619; Pred No. 0.00046;
35; Indels
                                                                                                                         507 others
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0; Gaps
0;
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RESULT 13
AX323765
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 TTTGGTTTAGATGTTTTTTG 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               40 GTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTC 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 34.0%;
Local Similarity 68.8%;
Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Arc, Jr.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D., Buck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Burky, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gao, J., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Norway rat.
Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Olek,A., Piepenbrock,C. and Berlin,K.
Diagnosis of diseases associated with dna transcription
Patent: WO 0192565-A 253 06-DEC-2001;
Epigenomics AG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AX323765.1 GI:18094513
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Ax323765
                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HTG; HTGS_PHASE1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AC126842.1 GI:21722906
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rattus norvegicus clone CH230-10C10,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note-"chemically treated genomic DNA (Homo sapiens)" 176 c 1712 g 2422 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic construct"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             185476 bp
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10, *** SEQUENCING IN PROGRESS
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Homsi,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E.,
Jacobson,B., Jia,Y., Johnson,R., Jolivet,S., Joudah,S.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Karlsson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C.,
Kratovic,J., Kureshi,A., Landry,N., Leal,B., Lewis,L.C., Lewis,L.,
Li,J., Li,Z., Lichtarge,O., Lieu,C., Liu,J., Liu,W., Louiseged,H.,
Lozado,R.J., Lu,X., Lucier,A., Lucier,R., Luna,R., Ma,J.,
Lozado,R.J., Lu,X., Lucier,A., Lucier,R., Luna,R., Ma,J.,
Maheshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E.,
Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M.,
Miner,G., Miner,Z., Mitchell,T., Mohabbat,K., Morgan,M., Mortis,S.,
Moser,M., Neal,D., Newtson,J., Newtson,N., Nguyen,A., Nguy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (10-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             * (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html)
* NOTE: This is a 'working draft' sequence. It currently
* consists of 69 contigs. The true order of the pieces
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          be preserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This record will be updated with the finished sequence as soon as it is available and the accession number will
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Grand-Pierre, N., Grant, G., Hagos, B., Heaford, A., Horton, L., Howland, J.C., Illev, I., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Locke, K., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McGurk, A., McKernan, K., McPheeters, R., McCarthy, M., McEwan, P., McGurk, A., McKernan, K., McPheeters, R., Meldrim, J., Meneus, L., Mihova, T., Miranda, C., Mlenga, V., Morrow, J., Murphy, T., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Olivar, T.M., Oliver, J., Peterson, K., Pierre, N.,
                                                                                                                                                                                                                                                         Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Boguslavo, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Domino, M., Doyle, M., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Grand, Blore, S., Goyette, M., Graham, L., Grand, B., Gran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 154696)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
Homo sapiens, clone RP11-216B13
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Homo sapiens clone RP11-216B13, WORKING DRAFT SEQUENCE, 35
unoxidered pieces.
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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JOURNAL

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Pisani,C., Pollara,V., Raymond,C., Riley,R., Rogov,P., Rothman,D., Roy,A., Santos,R., Schauer,S., Severy,P., Spencer,B., Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J., Tirrell,A., Travers,M., Trigillo,J., Vassiliev,H., Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Young,C., Zainoun,J., Zimmer,A. and Zody,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (12-MAR-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On May 28, 2000 this sequence version replaced gi:7657088.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          consists of 35 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOTE: This is a 'working draft' sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0,960731
Consensus quality: 135951 bases at least Q40
Consensus quality: 145629 bases at least Q30
Consensus quality: 148984 bases at least Q20
Insert size: 149000; agazose-fp
Insert size: 15196; sum-of-contigs
Quality coverage: 3.7 in Q20 bases; sum-of-contigs
Quality coverage: 3.7 in Q20 bases; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: sequence_submissions@genome.wi.mit.edu
------ Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequencing vector: M13; M77815;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Center project name: L580
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2: gap of 4777; cr
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               24050: gap of 100 b
26881: contig of 2831
26981: gap of 100 b
29772: contig of 2791
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3505: cor
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  29872: gap of
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    Genome Center

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                                                                                                                                                                                                                                                                                                                                                                                                      8097: contig of 1363 bp in length
37: gap of 100 bp
10306: contig of 2109 bp in length
                                                                                                          23950
                                                                                                                                              19608: contig of 1785 bp in length
08: gap of 100 bp
22007: contig of 2299 bp in length
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contig of 1843 bp in length
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contig of 1159 bp in length
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f 2791 bp in length
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57772 57871:
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62063 65728: contig of 3666 bp in length
65729 65828: gap of 100 bp
65829 69752: contig of 3924 bp in length
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32607: gap of 100 bp
36314: contig of 3707 bp in length
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104501: contig of 12538 bp in length
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84995; contig of 8408 bp in length
95; gap of 100 bp
91863; contig of 6768 bp in length
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61962: contig of 4091 bp in length
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48483: contig of 3443 bp in length
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76487: contig of 6635 bp in length
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Search completed: November 21, 2002, 03:10:02 Job time : 3494 \text{ secs}
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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AAK69569 AAK79238 AAS74886	AAS94925 AAL04302	ABK35928	ABV25399	ABK72318	ABK72317	ABK72316	ABK72314	ABK72313	ABK72312	ABK72310	ABK72309	ABK72308	AAC84595	AAN94725	AAN95403	AAN91363	AAD02543	ABK37512	AAS42222	ABQ56063	AAS84770	ABQ54676	AAH98115	ABQ88179	AAZ61522	AAH22143	AAC74939	ABN59848	ABL97872	ABL97871	AAL04979	AAL04978
Human immune/haema Human immune/haema DNA encoding novel	Human DNA sequence	sequence	0			Lymphona associate	Human C-myc prote1	<	DNA sequence encod	DNA sequence encod	Human SNORF40 rece	DNA encoding G-cou	Human cDNA encodin	Human ovarian anti	DNA encoding novel	Human ovarian anti	Murine 7-transmemb	Human osteoblast d	Cosmid CVO14 conta	Human chromosome 1	Human ORFX ORF494	Novel human coding	Human testicular a	Human testicular a	Human reproductive	Human reproductive						

## ALIGNMENTS

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RESULT 1
AAF30037
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                                                                                                                      Key
                                                                                                                                                   Apolipoprotein A-IV-related protein; AA4RP; human; biallelic marker; lipid metabolism; liver related disorder; diagnosis; gene therapy; chromosome 11; diabetes; obesity; coronary heart disease; ds.
                                                                                                                                                                                              Apolipoprotein A-IV-related protein gene (alternative sequence).
                                                                                                                                                                                                                                AAF30037;
                                                                                                                                                                                                                                                AAF30037 standard; DNA; 5381 BP.
                  intron
                                          exon
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                                                                                                             5'UTR
                                                                                                                                     Homo sapiens.
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931..1441
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919..930
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1499..1612
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Duclert A;

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 21-JUN-2000; 2000WO-IB01011
                                  04-JAN-2001.
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/number- :
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1725..2242
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base C or T"
replace(1241.T)
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/note= "17-42-319 probe"
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                                                                                            "17-41-250 probe"
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                                                                                                                                            RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                          Matches 121;
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20-DEC-1999;
21-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of an alternative genomic sequence (see also AAF30035) of human apolipoprotein A-IV-related protein (AA4RP). It encodes a 366-amino acid protein (see AAB20103). AA4RP is differentially expressed in obese mouse models, indicating a role in lipid metabolism related disorders. It appears to be the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding apolipoprotein A-IV-related proteins (AA4RP) and biallelic markers of AA4RP, useful for diagnosing lipid metabolism related disorders and/or liver related disorders -
            Human apolipoprotein A-IV-related protein (AA4RP) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 242-244; 260pp; English.
                                          23-APR-2001 (first entry)
                                                                                                          AAF30035 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5381 BP; 1104 A; 1434 C; 1709 G; 1132 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       probes and primers.
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99WO-1B02058
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DB 22; 0;

Length 5381;

Indels

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Gaps

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5'UTR
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12959..13469

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                                                                   "primer 20-828.rp"
                                                                                                                    "biallelic marker 20-853-415, polymorphic base C or T"
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as
"20-842-115 probe"
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A-IV-related protein (AA4RP) gene on chromosome 11. The gene
encodes a 366-amino acid protein (see AAB20103). AA4RP is

differentially expressed in obese mouse models, indicating a role
in lipid metabolism related disorders. It appears to be the human
homologue of rat regeneration associated protein (RAP3), which is
believed to be involved in liver regeneration, and is likely to
have a similar function. It also shows 52% similarity to
apolipoprotein A-IV, and is likely to have a similar function. The
invention also provides AA4RP cDNAs (see AAF30036) and polypeptides,
biallelic markers identified in the AA4RP gene and from genomic
regions flanking the gene, and methods for genotyping a nucleic
acid containing 1 or more of the biallelic markers. Also provided
is a method for detecting a statistical correlation between a
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                     12340 CAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCTTGC 12399
                                                                                                                                                                                                                                                                                                                                         12280 GCATTTGGGCTTGCTCAGAGGCCCTGCGAGTTGAGCTTTTCCTCATGGGG 12339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding apolipoprotein A-IV-related proteins (AA4RP) and biallelic markers of AA4RP, useful for diagnosing lipid metabolism related disorders and/or liver related disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JAN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 216-238; 260pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-1999;
21-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUN-2000; 2000WO-IB01011
 DNA transcription associated complementary genomic DNA #127.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               marker haplotype and/or between a biallelic marker haplotype and a phenotype. Diagnostic methods are provided for determining whether an individual is at risk of developing a lipid metabolism related disorder and/or a liver related disorder,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB; AAB20103.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-071485/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yen F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-JUN-1999;
                                23-APR-2002 (first entry)
                                                                       ABK28380;
                                                                                                          ABK28380 standard; DNA; 5377 BP
                                                                                                                                                                                                                                 121 C 121
                                                                                                                                                                                                                                                                                                       61
                                                                                                                                                                                                                                                                                                                                                                         1 GCATTTGGGCTTGCTCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTTCCTCATGGGG 60
                                                                                                                                                                                                                                                                                                       CAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCTTGC 120
                                                                                                                                                                                                                                                                                                                                                                                                               121;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Denison B,
                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-IB02058
99US-0469099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "20-853-415 probe"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bour B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 99.78;
                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                       Score 120.6; Db 44,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bíhain B,
                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 81001;
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DNA transcription associated gene; peptide nucleic acid; PNA-oligomer; PNA; cytosine methylation state; SNP; retroviral infection; gene; ds; single nucleotide polymorphism; adenosine deaminase deficiency; cancer; viral infection; Sezary syndrome; haematological disorder; tuberculosis; immunological disorder; Werner syndrome; developmental disorder; immunological disorder; werner syndrome; developmental disorder;
The invention relates to a nucleic acid, which comprises a segment of the chemically pretreated DNA of genes associated with DNA transcription from one of 346 sequences, and an oligomer, in particular an oligomucleotide or peptide nucleic acid (PWA)-oligomer that hybridises to or is identical to the chemically pretreated DNA of genes associated with DNA transcription. The set of oligomer probes are useful for detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acids or oligomers, useful for diagnosing or treating diseases associated with DNA transcription, e.g. immunological disorders, Werner syndrome, psoriasis, myocardial infarction, solid tumours or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-APR-2000; 2000DE-1019058
07-APR-2000; 2000DE-1019173
30-JUN-2000; 2000DE-1032529
01-SEP-2000; 2000DE-1043826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neurodegenerative disorder; Waardenburg syndrome; Niemann-Pick disease; myelodysplastic syndrome; myocardial infarction; hypertension; arthritis; angiogenesis; congenital heart disease; HDR syndrome; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-APR-2001; 2001WO-EP03973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID No 254; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPIG-) EPIGENOMICS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200192565-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            psoriasis; Rieger's syndrome; neurological disorder; erythropoiesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Piepenbrock C,
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cytosine methylation state and/or single nucleotide polymorphisms (SNPs) in a chemically pretreated genomic DNA. The nucleic acids are useful for diagnosing or treating diseases associated with DNA transcription (particularly with the methylation status), e.g. adenosine deaminase deficiency, viral infection, retroviral infection, Sezary syndrome, haematological disorders, immunological disorders, Werner syndrome, tuberculosis, developmental disorders, psoriasis, Rieger's syndrome, neurological disorders, neurodegenerative disorders, Waardenburg syndrome, Niemann-Pick disease, myelodysplastic syndrome, myocardial infarction, hypertension, anglogenesis, erythropolesis, congenital heart disease, HDR syndrome, arthritis, polyglutamine disorders, solid tumours or cancer. Sequences ABK28127-ABK28472 represent DNA transcription associated genomic DNA molecules of the invention. Note: The sequence data for this patent did not form part of the printed European Patent Office. specification but was obtained in electronic format directly from

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5376 TTCAACTTTTCCTCATAAAACAAATCTCACTTTCGCTCCAATTCCCTAAAACTCAAAATCC 5317

41 TTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCC 100

Matches

Local

Similarity

44.18; 77.88;

Score 53.4; DB 24; Pred. No. 3.1e-07; Mismatches

0;

Gaps

0;

Conservative

Sequence 5377 BP; 1117 A; 176 C; 1481 G; 2603 T;

0 other; Length 5377; Indels

δÃ

101 CTGGCCCAGATGCCTCTTGCC 121

5316 CTAACCCAAATACCTCTTACC 5296

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RESULT 4
ABK28379
The invention relates to a nucleic acid, which comprises a segment of the Cohemically pretreated DNA of genes associated with DNA transcription from the cone of 346 sequences, and an oligomer, in particular an oligonuclectide one of 346 sequences, and an oligomer, in particular an oligonuclectide or peptide nucleic acid (PNA)-oligomer that hybridises to or is identical to the chemically pretreated DNA of genes associated with DNA construction. The set of oligomer probes are useful for detecting the cytosine methylation state and/or single nucleotide polymorphisms (SNPs) in a chemically pretreated genomic DNA. The nucleic acids are useful for Cd diagnosing or treating diseases associated with DNA transcription concept of the methylation status), e.g. adenosine deaminase deficiency, viral infection, retroviral infection, Sexary syndrome, tuberculosis, developmental disorders, psoriasis, Riegar's syndrome, conversed to the cological disorders, neurodegenerative disorders, Warner syndrome, neurological disorders, myelodysplastic syndrome, mycoardial infarction, hypertension, angiogenerative disorders, wardenburg conversed to the syndrome, arthritis, polyglutamine disorders, solid tumours conversed to DNA molecules of the invention.

CC associated genomic DNA molecules of the invention of the printed specification but was obtained in electronic format directly from the culture of the printed content of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     viral infection; Sezary syndrome; haematological disorder; tuberculosis;
immunological disorder; Werner syndrome; developmental disorder;
psoriasis; Rieger's syndrome; neurological disorder; erythropolesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acids or oligomers, useful for diagnosing or treating diseases associated with DNA transcription, e.g. immunological disorders, Werner syndrome, psoriasis, myocardial infarction, solid tumours or cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-APR-2000; 2000DE-1019058.
07-APR-2000; 2000DE-1019173.
30-JUN-2000; 2000DE-1032529.
01-SEP-2000; 2000DE-1043826.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neurodegenerative disorder; Waardenburg syndrome; Niemann-Pick disease; myclodysplastic syndrome; myocardial infarction; hypertension; arthritis; angiogenesis; congenital heart disease; HDR syndrome; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA transcription associated gene; peptide nucleic acid; PNA-oligomer; PNA; cytosine methylation state; SNP; retroviral infection; gene; ds; single nucleotide polymorphism; adenosine deaminase deficiency; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA transcription
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID No 253; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-APR-2001; 2001WO-EP03973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polyglutamine disorder; solid tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-APR-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Piepenbrock
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RESULT 5
AAD26473/c
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                                                                                                                                                                                                                                                                                                      23-JUN-2000;
30-JUN-2000;
07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immune disorder; atherosclerosis; Crohn's disease; Hodgkin's disease; Acquired Immune Deficiency Syndrome; AIDS; Addison's disease; anaemia; allergy; asthma; adult respiratory distress syndrome; multiple sclerosis; autoimmune thyroiditis; bronchitis; diabetes mellitus; osteoarthritis; Good pasture's syndrome; Graves' disease; pancreatitis; psoriasis; rheumatoid arthritis; ulcerative colitis; citrhosis; Cushing's syndrome; hepatitis; hypothyroidism; cerebral palsy; cataract; angina pectoris; cardiovascular disease; hypertension; vasculitis; myocarditis; obesity; congestive heart failure; ischaemic heart disease; lung tumour; gout;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5377 BP; 1067 A; 176 C; 1712 G; 2422 T; 0 other;
           New polypeptides, useful for diagnosing, treating or preventing disorders of growth and development, cardiovascular and lipid, diseases such as cancer, comprise human kinase polypeptides
                                                                                                                       Yue H, Lal P, Bandman O, Borowsky ML, Au-Young J, Lu Y; Gandhi AR, Tribouley CM, Walia NK, Yao MG, Lu DAM, Green Ramkumar J, Griffin JA, Kearney L; Burford N, Nguyen DB, Baughn MR, He A, Thornton M, Hafalia A, Patterson C, Gur Lo TP, Khan F, Recipon SA, Azimzai Y, Policky JL, Ding L Grether M, Elliott VS, Thangavelu K, Batra S, Ison CH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; kinase; PKIN-26; cancer; leukaemia; adenocarcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human kinase PKIN-26 cDNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                        14-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                      20-DEC-2001.
                                                                                                                                                                                                                                        (INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                     15-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          40 GTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTC 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GTTTAGTTTTTTTTATGGGGTAAATTTTATTTTCGTTTTAGTTTTTGGGGGTTTAGAGTT 60
                                                                                            2002-090207/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     liver; Niemann-Pick's disease; gene therapy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                              AAE16280.
                                                                                                                                                                                                                                                                                        2000US-212073P.
2000US-213467P.
2000US-215651P.
2000US-216605P.
2000US-218372P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                        2001WO-US19444
                                                                                                                                                                                                                                                                          2000US-228056P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product- "Human PKIN-26 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 41.2; DB 2
Pred. No. 0.0025;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24;
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                                                                                                                           Ding L;
CH;
                                                                                                                                                                                        Lu Y;
Greenwald SR;
                                                                                                                                                           Gururajan R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                osteoporosis;
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AA160549/c
ID AA1605
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                                                                                                                                                 21-JAN-2000;
25-APR-2000;
09-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; nootropic; immunosuppressant; cytostatic; gene therapy; cance peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; heemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                          03-AUG-2000;
                                                                                                                                                                                                                                                                                                      26-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200153312-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-OCT-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAI60549 standard; cDNA; 3372 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2212 ACACAATCTCTCCCGTCCCTGGAGGCCAGCTCCCCCG 2176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2272 GTGCTACCTGGTCTCCCAAGGCCCTTCAGACTTTTCCACGTTGGTCTCGTGTGGCACATG 2213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to human kinase PKIN proteins and their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
   14-SEP-2000;
19-OCT-2000;
                                                                                                                 19-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                 26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human polynucleotide SEQ ID NO 4538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGGCTTGCTCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATC 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PKIN-26 CDNA.
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                                       2000US-0620312.
2000US-0653450.
2000US-0662191.
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                                                                                                                                                                                                                                                                                                      2000WO-US34263
2000US-0693036
                                                                                                                                                 2000US-0552317.
2000US-0598042.
                                                                                                                                                                                                                           2000US-0488725.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        197; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     553 A; 678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; 812 G; 416 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 2459;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                     AA158763/c
                                                                                                                                                                                                                                                                                                                                                                  RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tang
Wang
           26-DEC-2000; 2000WO-US34263.
                                                                                                                                                 chemokinetic;
                                                                                                                                                             peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                                                                                                                                                                                                                                                                                                             2187 ACACAATCTCTCCCGTCCCTGGAGGCCAGCTCCCCCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3372 BP; 734 A; 975 C; 1071 G; 592 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C.N.S disorders. Note: The sequence data for this pattent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 4538; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel nucleic acids and such as central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-NOV-2000; 2000US-0727344
                                                                         WO200153312-A1
                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                            Human polynucleotide SEQ ID NO 966
                                                                                                                                                                                                                                                                                                                                     AAI58763 standard; cDNA; 3446 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       assays for receptor activity, arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-442253/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HYSE-) HYSEQ INC
                                                                                                                                  leukaemia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polypucleotides are useful in gene therapy. A composition containing a polypeptide or polypucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                         67 TYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTG 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 GGGCTTGCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATC 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Joca 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTGCTACCTGGTCTCCCAAGGCCCTTCAGACTTTTCCACGTTGGTCTCGTGTGGCACATG 2188
                                                                                                                                                                                            nootropic; immunosuppressant; cytostatic; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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Wang Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                  thrombolytic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Asundi V, Ch
Wehrman T, X
Goodrich R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wehrman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptides, useful
system injuries -
                                                                                                                                               drug screening; arthritis; inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 29.4;
Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drmanac RT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Qian XB,
Yang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for treating disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ren F, 1
Zhang J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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AAK70514/c
ID AAK70514 standard; DNA; 12730 BP.
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   PD XXX XXX DT XXX
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-SEP-2000;
19-OCT-2000;
29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activity, chemotactic/chemokinetic activity, hemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tang
                                                                                                                                                                                                                                                                                                                                                                                                                           2319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel nucleic acids and polypeptides, useful such as central nervous system injuries -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang
                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 966; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-442253/47.
P-PSDB; AAM39607.
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   09-AUG-2001
                                   WO200157182-A2
                                                                                                                                                                                              06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3446 BP; 759 A; 991 C; 1094 G; 602 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C.N.S disorders.

Note: The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhao
                                                                                                     cytostatic; gene therapy; vaccine; metastasis; ds.
                                                                                                                      Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                         Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25326.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specification.
                                                                                                                                                                                                                                AAK70514;
                                                                                                                                                                                                                                                                                                                                                                                       67
                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 GGGCTTGCTCCTCAGAGGCCCCTGCGAGTGGAGTTCAGCTTTTTCCTCATGGGGCAAATC 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QΑ,
                                                                    sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YT,
                                                                                                                                                                                                                                                                                                                                                      ACACAATCTCCCCGTCCCTGGAGGCCAGCTCCCCCG 2223
                                                                                                                                                                                                                                                                                                                                                                       TYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTG 103
                                                                                                                                                                                                                                                                                                                                                                                                                         GTGCTACCTGGTCTCCCAAGGCCCTTCAGACTTTTCCACGTTGGTCTCGTGTGGCACATG 2260
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Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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2000US-0552317.

2000US-05598042.

2000US-06620312.

2000US-0662315.

2000US-0662191.

2000US-0593036.

2000US-0593036.
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                                                                                                                                                                                            (first entry)
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Wehrman T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goodrich R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24.3%;
55.7%;
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, Xu C, Xue AJ,
R, Drmanac RT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 29.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
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Yang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhang J;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3446;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
05-SEP-2000;
05-SEP-2000;
                                                                                                                                                                                                                                                                              08-SEP-2000;
12-SEP-2000;
14-SEP-2000;
14-SEP-2000;
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08-SEP-2000;
08-SEP-2000;
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06-SEP-2000;
08-SEP-2000;
                                                                                                                  14-SEP-2000;
21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
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14-SEP-2000;
14-SEP-2000;
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07-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-AUG-2000;
22-AUG-2000;
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                                                                                                                                                                                                                                                               14-SEP-2000;
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                           2000US-0234998.
2000US-0235484.
2000US-0235834.
2000US-0235836.
2000US-0236327.
                                                                                                                                                                                                                                                                                               2000US - 0229343

2000US - 0229344

2000US - 0229345

2000US - 0229509

2000US - 0229513

2000US - 0230437

2000US - 0231242

2000US - 0231243

2000US - 0231244

2000US - 0231244

2000US - 0231244

2000US - 0231240

2000US - 0231414

2000US - 0231414

2000US - 0231414

2000US - 0232397
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2000US -0225267
2000US -0225268
2000US -0225270
2000US -0225447
2000US -0225757
2000US -0225758
2000US -0225759
2000US -0225759
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2000US-0224519.
2000US-0225213.
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2000US-0232399.
2000US-0232400.
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13-OCT-2000;
20-OCT-2000;
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2000US-0251868.
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RESULT 9
AAX99032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid sequences given in AAM82170 to AAM91921. (1) have cytostatic cartivity, and can be used in gene therapy and vaccine production. (1) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (1) expression. For cexample, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (1) by expressing inactive proteins or to supplement the patients own production of (1) Additionally, (1) concerns and caids into a host cell and culturing the cell to express the protein. (1) proteins and polynucleotides may be used to prevent, and diagnose and treat immune/haematopoietic related diseases, especially cancers and cancer metastases of haematopoietic derived cells. AAK64703 to AAK87694 represent human immune/hacmatopoietic antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169
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Human validated cancer cell derived cDNA #354.
                                           24-SEP-1999
                                                                                  AAX99032;
                                                                                                                           AAX99032 standard; cDNA; 845 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure;
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                                                                                                                                                                                                                                                                                                                                         GGGCTTGCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATC 66
                                                                                                                                                                                                                                                      TYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTG
                                                                                                                                                                                                                                                                                                             GTGCTACCTGGTCTCCCAAGGCCCTTCAGACTTTTCCACGTTGGTCTCGTGTGGCACATG 1940
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                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                24.3%;
55.7%;
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 12730;
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21-DEC-1998;
23-DEC-1997;
03-APR-1998;
21-OCT-1998;
27-OCT-1998;
                                                                                                                                                                                                                                                               Cancer; human; colon; breast; lung; transmembrane receptor; ATPase; integral membrane protein; aspartyl protease; GATA family, wnt family: transcription factor; G-protein alpha subunit; protein phosphatase; phorbolester binding protein; diacylylycerol binding protein; trypsin; protein kinase; tyrosine phosphatase; developmental signalling protein; protein kinase; tyrosine phosphatase; developmental signalling protein; wW/rsp5/WWP domain; therapy; forensic; genetic mapping; diagnostic; detection; treatment; cervical; melanoma; colorectal adenocarcinoma; wilm's tumour; retinoblastoma; sarcoma; myosarcoma; lung carcinoma; wilm's tumour; retinoblastoma; sarcoma; myosarcoma; lung carcinoma;
                                                                                                                                                                   W09933982-A2
                                                                                                    22-DEC-1998;
                                                                                                                                   08-JUL-1999
                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                    leukemia;
                                                                                                                                                                                                                                    lymphoma; dysplasia; hyperplasia; endometrium; adrenal; ss.
98US-0217471.
97US-0068755.
98US-0080664.
98US-0105234.
98US-0105877.
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Best Local :
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Lamson G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              agents and in forensics, genetic analysis, mapping and diagnostic applications. In particular, the product can be used for the detection and management of cancers. They can be used for treating e.g. cervical cancers, melanomas, colorectal adenocarcinomas, Wilm's tumour, sarcomas, retinoblastoma, myosarcomas, lung carcinomas, leukemias, such as chronic myelogenous leukemia, promyelocytic leukemia, monocytic leukemia, and myeloid leukemia, and lymphomas such as histiocytic lymphoma, anhydric hereditary ectodermal dysplasia, congenital alveolar dysplasia, epithelial dysplasia of the cervix, fibrous dysplasia of bone, and mammary dysplasia, hyperplasias, e.g. endometrial, adrenal, breast,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 555; 591pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stache-Crain B, Sudduth-Klinger J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                     prostate or thyroid hyperplasias or pseudoepitheliomatous hyperplasia the {\tt skin.}
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Human; reproductive system related antigen; reproductive system disorder;
                                Human reproductive system related antigen DNA SEQ ID NO: 7666.
                                                                                                                              AAL04978 standard; DNA; 1768 BP
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 845 BP; 215 A; 144 C; 170 G; 236 T; 80 other;
                                                              21-NOV-2001 (first entry)
                                                                                                AAL04978;
                                                                                                                                                                                                            134 AAAAACCTGCACACGCACCANTTCCCGTNNC 164
                                                                                                                                                                                                                                           61 CAMATCTYACTTTCGCTCCAGTTCCTGGGGC 91
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                                                                                                                                                                                                                                                                                                         1 GCATTTGGGCTTGCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGG 60
                                                                                                                                                                                                                                                                                                                                          49; Conservative
                                                                                                                                                                                                                                                                                                                                                         Similarity
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J, Garcia PD, Garcia V, Giese K,
Kassam A, Kennedy GC, Kita D, L.
Leshkowitz D, Pot D, Randazzo F,
                                                                                                                                                                                                                                                                                                                                                       24.1%; Score 29.2;
53.8%; Pred. No. 11;

    Mismatches

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Kita D, Labat I;
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33-AUG-2000
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01-SEP-2000
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05-SEP-2000
06-SEP-2000
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08-SEP-2000
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08-SEP-2000;
                        2000US-0232400.
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25-SEP-2000 26-SEP-2000 27-SEP-2000 27-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 02-OCT-2000 02-OCT-2000 02-OCT-2000 02-OCT-2000 01-OCT-2000 01-OCT-2000

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2000US-0250391.

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2000US-0251986.

2000US-0256719.

2000US-0251479.

2000US-0251856.

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Best Local
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04 - EB - 2000

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12 - MAR - 2000

16 - MAR - 2000

17 - MAR - 2000

18 - APR - 2000

19 - MAY - 2000

07 - JUN - 2000

28 - JUN - 2000

07 - JUL - 2000

07 - JUL - 2000

11 - JUL - 2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a
                                                                                                                                                                                                                                                                                                                                                                                       WO200155320-A2
                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; reproductive system related antigen; reproductive system disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human reproductive system related antigen DNA SEQ ID NO: 7667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAL04979;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1100 TGCTATTGTCAAATGTGCTGAAAGACAAGTTGTCCCTTTACCCCTTAGGAGAGAGGCTCCT 1159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1768 BP; 445 A; 440 C; 461 G; 422 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                        02-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12 TGCTCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTTCCTCATGGGGCAAATCCTYACT 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72 TTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAG 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Barash SC,
                              2000US-0189874
2000US-0199076
2000US-0199076
2000US-029515
2000US-0205515
2000US-0214886
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2000US-0216847
2000US-0216480
2000US-0216480
2000US-0217487
2000US-0217487
2000US-0217486
2000US-0217486
2000US-0217486
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                                                                                                                                                                                                                                                                                          2000US-0179065
                                                                                                                                                                                                                                          2000US-0186350
                                                                                                                                                                                                                                                          2000US-0184664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 28.8;
Pred. No. 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 22; Length 1768;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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06-DEC-2000
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14 - AUG

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2000US-0232398
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2000US-0234239
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2000US-0236871
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2000US-0227009.
2000US-0228924.
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Best Local S
Matches 54
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08-NOV-2000
17-NOV-2000
17-NOV
                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a protein of the invention.
                                     Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                       Sequence 1768 BP; 445 A; 441 C; 459 G; 422 T; 1 other;
                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-465570/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME
                                                                        12 TGCTCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACT 71
72 TTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAG
                                                                                                            1 Similarity 55.1 54; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Barash SC,
                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 7667; 1297pp + Sequence Listing; English.
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2000US-0251938.
2000US-0251938.
2000US-0251856.
2000US-0251858.
                                                                                                                             23.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ruben SM;
                                                                                                            Score 28.8; D: Pred. No. 18; 1; Mismatches
                                                                                                              ;;
                                                                                                                                                 DB 22;
                                                                                                              43;
                                                                                                                                                 Length 1768;
                                                                                                              Indels
                                                                                                              0,
                                                                                                            Gaps
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114 - AUG118 - AUG22 - AUG22 - AUG22 - AUG22 - AUG23 - AUG23 - AUG23 - AUG30 - AUG310 - SEP01 - SEP01 - SEP01 - SEP05 - SEP05 - SEP06 - SEP06 - SEP08 - SEP-

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23-AUG-2000

23-AUG-2000

01-SEP-2000

01-SE
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07-JUN 2000
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07-JUL 2000
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04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; testicular antigen; testes; cancer; metastasis; immune disorder; reproductive system disorder; urinary system disorder; gene therapy; cardiovascular disorder; respiratory disorder; neurological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JAN-2001;
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2000US-019874
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2000US-0198123
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2000US-0184664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO: 2523
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2000US-0232401.
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2000US-0246532

2000US-0246639

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2000US - 0236802
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2000US-0235484.
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2000US-0235836.
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2000US-0236327.
2000US-0236367.
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2000US-02464777.
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2000US-0231413
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2000US-0233064
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2000US-0241787.
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2000US-0246523.
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2000US-0241
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RESULT 13
ABL97872
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Matches 54
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08-DEC-2000;
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08-DEC-2000;
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17-NOV-2000;
01-DEC-2000;
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05-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides the protein and coding sequences of 973 human testicular antigens, and fragments of their genomic sequences. The sequences can be used in the treatment of cardiovascular, uninary system, reproductive system, immune, respiratory, neurological and gastrointestinal disorders, infections, and particularly cancer, especially testicular cancers. The present sequence is a DNA encoding a
                                                                                                                          Human; testicular antigen; testes; cancer; metastasis; immune disorder; reproductive system disorder; urinary system disorder; gene therapy; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disease; infection; cytostatic; gene; ds.
                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                            ABL97872 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                  1160 TACACTGACATTCAAAGTCCTCAATGACCCGGCCCCAG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 2523; 766pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding 973 human testicular antigen polypeptides, useful for preventing, diagnosing and/or treating testicular cancer
                                                        WO200155317-A2
                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                    21-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                          ABL97872;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein fragment of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAG 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TGCTCTCCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACT 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-483232/52.
                                                                                                                                                                                                                    testicular antigen encoding DNA fragment SEQ ID NO: 2524.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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2000US-0251999.
2000US-0254097.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 28.8; DI
Pred. No. 18;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 1768;
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Nucleic acids encoding 973 human testicular antigen polypeptides
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               WPI; 2001-483232/52.
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                                          (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences of 973 human testicular antigens, and fragments of their genomic sequences. The sequences can be used in the treatment of cardiovascular, urinary system,
             The present invention provides the protein and coding sequences of 444 novel human proteins. These were isolated from expressed sequences tags (ESTS). They can be used to stimulate cell growth, to regulate haematopoiesis e.g. to treat aplastic anaemia, to help tissue regrowth e.g. In burn treatment, to regulate the immune system e.g. to treat multiple sclerosis, to regulate activin or inhibin e.g. to treat infertility, to regulate haemostasis or thrombolysis e.g. to treat
                                                                                                                                                  Claim 1; SEQ ID NO 259; 509pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective; antiparkinsonian; protein therapy; EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antianaemic; vulnerary; antiinflammatory; immunomodulator; antiinfertility; cerebroprotective; cytostatic; rheumatic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN59848 standard; cDNA; 3549 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein fragment of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gastrointestinal disorders, infections, and particularly cancer, especially testicular cancers. The present sequence is a DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; SEQ ID NO 2524; 766pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for preventing, diagnosing and/or treating testicular cancer
                                                                                                                                                                                    encoded polypeptide such as cancer and multiple sclerosis
                                                                                                                                                                                                                                                    WPI; 2002-292408/33.
                                                                                                                                                                                                                                                                                                  Tang YT,
                                                                                                                                                                                                                                                                                                                                                                    11-SEP-2000; 2000US-0659671
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expressed sequence tag; gene; ss.
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                                                                                                                                                                                                  An isolated polynucleotide for treating diseases associated with
                                                                                                                                                                                                                                     P-PSDB; ABB97435
                                                                                                                                                                                                                                                                                    Xue AJ,
                                                                                                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC
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                                                                                                                                                                                                                                                                                                    Liu
   cancer, to
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                                                                                                                                                                                                                                                                                                  Zhou P,
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Pred. No. 18
   IOI
                                                                                                                                                                                                                                                                                  Drmanac RT;
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drugs,
 to treat inflammatory conditions
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulncrary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antianaemic; gene therapy; cancer; proliferative disorder; hypotrension; neurodegenerative disorder; osteoarthritis; graft vs host disease; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoslobinutia; burn; wound; constitution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. rheumatoid arthritis, and to treat nervous system disorders e.g. Parkinson's disease. The present sequence is a coding sequence of the
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AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human ORFX open reading frames 1 to 3161. The ORFX sequences have activities such as: cytostatic; hepatotropic; vulnerary;
                                                                            Claim 5; Page 962-964; 5507pp; English.
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                                                                                                                                                                                                                                                                                                                                                       02-APR-1999;
05-APR-1999;
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                                                                                                                     neurodegenerative disorders and cardiovascular disease
                                                                                                                                       Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders,
                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                      30-MAR-2000; 2000US-0540763
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thrombosis; contraceptive; ss.
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DB; AAB40730.
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99US-0127636.
99US-0127728.
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Best Local :
                                                                                                         1548 GTTCCTGCACCACCTGC 1564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bacterial or fungal infection, malaria, autoimmune disorders, asthma, allergies, aplastic anaemia, burns, wounds, bone and cartilage damage, nocturnal haemoglobinuria, antiinflammatory disease; to enhance coagulation; to inhibit thrombosis; and as a contraceptive.
                                                                                                                                                                                                               1488 GGATGTCAGCCTGACCTCGCCTGGCCAAACTCCCCAGTGAGACCATTTTCGTGGGCTGCGA 1547
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4224 BP; 961 A; 1150 C; 1123 G; 988 T; 2 other;
                                                                                                                                                                                                                                                              37 GGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCCTCAGA 96
                                                                                                                                                           97 GTCCCTGGCCCAGATGC 113
                                                                                                                                                                                                                                                                                                                46;
                                                                                                                                                                                                                                                                                                                                               Similarity
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    Mismatches

                                                                                                                                                                                                                                                                                                                                            Score 28.6;
Pred. No. 2
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

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OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Minimum DB seq length: 0
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121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16154066 seqs, 8097743376 residues
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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33.2	33.2	33.2	33.2	33.2	33.8	Score
27.4	27.4	27.4	27.4	27.4	27.9	Query Match 1
698	692	573	457	448	285	Query Match Length DB
10	10	10	3	12	13	: <del>B</del>
AW793634	BE256387	BE252183	BG982886	BF724998	BI004187	esult Query NO. Score Match Length DB ID
AW793634 MR1-UM000	BE256387 601117848	BE252183 601113780	BG982886 PM0-CN015	BF724998 bx10h11.y	BI004187 MR4-HN006	Description

0 0 40 0 0 41 0 0 42 0 0 44 4 43	C 33 34 37 38	c 27 c 27 c 29 c 31	c 23 c 23 c 23 c 23	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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A1746820 A1182101 BF659555 B1221371 B1554370 B1144788 BF532593	50031 50031 E6267 E6321 E6321 B4785 B1088 B1088	BB318044 AL520680 AL559313 BM470206 AZ856127 AZ856127 BB926542	AZZABBO ALSZ6B99 AZZ75B14 AZG11195 BH095750 ALSZ9993 AK013574 AQ663957	BG744224 AL528119 BM019314 B0896139 AL560785 AL560022 AL560022 AL56053 AL583065 BG762959 BG762959
A1746820 u107912 y A1746820 u1079609 y A182101 ud73e09 y BF65955 maa32f08. B1221371 602938928 B1554370 603235872 B114478B 602910003 BF532593 602074649	vg07 8 uul 4 uul 1 BB4 2 Mus	BB31804 .L520680 .L559313 .AGENCOU 2M01601 .L526542	L526899 L526899 RPCI-23- 1M0436N0 RPCI-24- L529993 L529993 Mus musc HS_2136_	BG744224 602723330 AL528119 AL528119 BM019314 603647443 BQ096139 AGENCOURT AL520679 AL526079 AL560725 AL560022 AL560022 AL560023 AL56053 AL56053 AL56053 AL583065 BG762959 602735012 BF795030 602256150 A7378865 1M0552P17

## ALIGNMENTS

Locus

KEYWORDS SOURCE ORGANISM DEFINITION ACCESSION RESULT 1 BI004187/c COMMENT REFERENCE VERSION JOURNAL MEDLINE TITLE AUTHORS Dias Neto,E., García Correa,R., Verjovski-Almeida,S., Briones,M.R., Ragai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and 285 bp mrNA linear EST 13-JUN-200 MR4-HN0065-210201-004-a05 HN0065 Homo sapiens cDNA, mrNA sequence. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 285) Tel: +55-11-2704922 Fax: +55-11-2707001 Shotgun sequencing of the human transcriptome with ORF expressed sequence tags Homo sapiens BI004187.1 Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP, Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research Contact: Simpson A.J.G. 20202663 Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000) GI:14408261 EST 13-JUN-2001

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RESULT 2
BF724998/c
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR4&t2=MR4-HN0065-
210201-004-a05&t3-2001-02-21&t4=1)
Seq primer: puc 18 forward
High quality sequence stop: 285.
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BF724998
                                                                                                                                                                                                                                                                      Plate: 10 row: h column: 11
Seq primer: M13RP1 reverse primer (ABI).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wistow, G.J., Bernstein, S., Behal, A. and Smith, D. NEIBANK: EST analysis and bioinformatics for ocular genomics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 448)
                                                                                                                                                                                                                                                                                                                                                                                                                 National Eye Institute
6/331, NIH, Bethesda, MD 20892-2740, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Section on Molecular Structure and Function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Wistow G
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                                                                                                                                                                                                                                                                                                                                                  graeme@helix.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note-*Organ: head_normal; Vector: pucl8; Site_1: SmaI; Site_2: SmaI, A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196.716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

1 others
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/note="Organ: Eye; Vector: pCMVSPORT6; Post-mortem iris tissue was pooled from 10 individuals ranging in age from
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                                                                                           /tissue_type="Iris"
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/db_xref="taxon:9606"
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                                                                      /dev_stage="Adult"
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                                               /lab_host="EMDH10B"
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В Š 뭥

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                                                                                                                                                                                                                                        This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl-pMO&t2-PMO-CN0155-050301-003-f11&t3-2001-03-05&t4-1)
Seq.primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
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                                                                                                                                                                                                                    High quality
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence tags
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Simpson, A.J.
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                                                                                                                                                                                                                                                                                                                                                                               Email: asimpson@ludwig.org.br
                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: +55-11-2704922
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/note-*Organ: colon_normal; Vector: pucl8; Site_1: SmaI;
Site_2: SmaI; A mini-library was made by cloning products
                                                      /dev_stage="Adult"
                                                                             /clone_lib="CN0155"
                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                             Pocari
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455 TGGGGGTCTTCTTTCTTCGTTCAAATCCAACTTCCTCTTCCCTTTGGGAGACTCCT 396
                       36 TGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAG 95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90 TGGGGGTCTTCTTTCTTCTTGGTTCAAATCCAACTTCCTCTTCCCTTTGGGAGACTCCT 149
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BE252183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                         Conservative
                                                                                                                                                                                                     /tissue_type="retinoblastoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by Oligo-dT priming. Directionally
cloned into ECORL/XhoI sites using the following 5;
adaptor: GGCACGAG(G). Library constructed by Ling Hong
in the laboratory of Garald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MGC Library."
34 a 193 c 146 g 100 t
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:3354499"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="NIH_MGC_16"
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Homo sapiens
                                                                                                                                                      MR1-UM0008-290200-002-c06 UM0008 Homo sapiens cDNA, mRNA sequence.
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Tissue Procurement: ATCC
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National Institutes of Health, Mammalian Gene Collection (MGC)
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/tissue_type="refinoblastoma"
/lab host="9H10B (phage-resistant)"
/lab host="9H10B (phage-resistant)"
/note="0rgan: eye; Vector: pOTB); Site_1: XhoI; Site_2:
/note="0rgan: eye; Vector: priming. Directionally
/cloned into EcoRI/XhoI sites using the following 5'
/cloned into EcoRI/XhoI sites using the following 5'
/cloned into EcoRI/XhoI sites using the following 5'
/cloned into EcoRI/XhoI sites using the following 15'
/cloned into EcoRI/XhoI sites using the following 15'
/cloned into EcoRI/XhoI sites using the following 15'
/cloned into EcoRI/XhoI sites using Tap-cona (university of california, Berkeley) using Zap-cona synthesis kit
/cloned into EcoRI/XhoI sites using Tap-cona synthesis kit
/cloned into EcoRI/XhoI sites using Tap-cona synthesis kit
/cloned into EcoRI/XhoI sites using the following 15'
/cloned into EcoRI/XhoI sites using the following 5'
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/clone="IMAGE:3358330"
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                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 787)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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High quality sequence stop: 696.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
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Ludwig Institute for Cancer Research
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Unpublished (1999)
                                                                                                                                                            Homo sapiens
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Tissue Procurement: Dr. Daniel McVicar, DBS/NCI
Tissue Procurement: Dr. Daniel McVicar, DBS/NCI
CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                              Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization Unpublished (2001)
                                                                                                                                                                                                                                            BP 191 91006 EVRY cedex - France
Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
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                                                                                                                                                                                                                                                                                                                         Contact: Genoscope
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http://image.llnl.gov
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//lab_host="pid108 (phage resistant)"
/note="organ: blood; Vector: poTPP; Site_1: XhoI: Site_2:
ECORI; cDNA made by oligo-dT priming. Directionally cloned
into ECORI/XhoI sites using the following 5 adaptor:
GCCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
NIH_MGC Library."
a 251 c 232 g 124 t
                                                                                            /clone="CSODC028YB17"
/clone_lib="LTI_NFL003_NBC3"
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/db_xref="taxon:9606"
/clone="IMAGE:4849686"
/clone_lib="NIH_MGC_106"
                                                                                                                                                /organism~"Homo sapiens"
/db_xref="taxon:9606"
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/note="Organ: brain; Vector: pCMVSPORT 6; lst strand cDNA
                       /tissue_type="neuroblastoma cells"
/lab_host="DH10B"
                                                                         /sex="male"
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Pred. No. 4.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLCM1897 row: e column: 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Clone distribution: MGC clone distribution information can be
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                                                                                                                                             /clone_ilb="NAGE.98"
/clone_ilb="NIH_MGC_98"
/clone_ilb="NIH_MGC_98"
/clone_ilb="NIH_MGC_98"
/tissue_type="sstrocytoma grade IV, cell line"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: brain; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: brain; Vector: poTB7; 
                                                                                                                                 Note: 272
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Pred. No. 4.2;
Score 33.2; DB 13; Length 832; Pred. No. 4.2;
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                                                           AL520679 LTI_NFL004_NBC2
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Tissue Procurement: ATCC
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                                      prime, mkNA sequence.
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/lab_host="DH10B (phage-resistant)"
/note="Organ: eye; Vector: pcMV-SPORT6; Site_1: Not1;
Site_2: Sall; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Library constructed by Life
Technologies."
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                                                                                                        BP 191 91006 EVRY cedex - France
Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
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/lab_host-"DH10B"
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/clone="CSODL004YA13"
                    /db_xref="taxon:9606"
                                         /organism="Homo sapiens"
                                                                                   Cocation/Qualifiers
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Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length CDNA libraries and normalization
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Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
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/tissue_type="B cells from Burkitt lymphoma"
/note="Vector: pcMvSPORT 6; Site_1: Not1; lst strand cDNA
/note="Vector: pcMvSPORT 6; Site_1: Not1; lst strand cDNA
/note="Vector: Disparsed with a Not1: oligo(dT) primer. Five prime end
enriched, double-stranded cDNA was digested with Not 1 and
cloned into the Not 1 and Eco RV sites of the pcMvSPORT 6
vector. Library was normalized. Library was constructed by
Life Technologies. Contact: Feng Liang Life Technologies,
a division of Invitrogen 9800 Medical Center Drive
Rockville, Maryland 20850, USA Fax: (1) 301 610 8371
Email: fliang@lifetech.com URL:
                                                                                                 note="Vector: pCMVSPORT 6; 1st strand cDNA was primed
with a NotI-oligo(dT) primer. Five prime end enriched,
double-stranded cDNA was digested with Not I and cloned
into the Not I and Eco RV sites of the pCMVSPORT 6 vector.
Library was constructed by Life Technologies, Contact :
Peng Llang Life Technologies, a division of Invitrogen
9800 Medical Center Drive Rockville, Maryland 20850, USA
Fax : (1) 301 610 8371 Email : filang@lifetech.com URL :
http://fulllength.invitrogen.com"
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/sex="male"
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/db_xref-"taxon:9606"
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/lab_host="DH10B"
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Score 33.2; DB 9; Length 984; Pred. No. 4.6;
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                                                                                                                                                                                                                                                                                                                                        RESULT 15
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   ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   726 GGGGCTTGGAGGTGCTGCCTATGGCC 701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      786 TGGGGGTCTTCTTTCTTCTTGGTTCAAATCCAACTTCCTCTTCCCTTTGGGAGACTCCT 727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AL556053 LTI_NFL006_PL2 Homo sapiens cDNA clone CS0DK010YL22 5
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BP 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
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/clone_lib="LTI_NFL006_PL2"
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Pred. No. 4.6;
0; Mismatches 33;
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TITLE
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                                                                                            760 GGGGCTTGGAGGTGCTGCCTATGGCC 735
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1 (bases 1 to 996)

Li,W.B., Gruber,C., Jessee,J. and Polayes,D.

Full-length cDNA libraries and normalization

Unpublished (2001)

Contact: Genoscope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: from Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="B cells from Hurkitt lymphoma"
/note="Vector: pCMVSPORT 6; Site_1: Not1; Ist strand cDNA
was primed with a Not1-oligo(dT) primer. Five prime end
enriched, double-stranded cDNA was digested with Not I and
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Copyright (c) 1993 - 2002 Compugen Ltd.
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/cgn2_6/ptodata/l/ina/5B_COMB.seq:*
/cgn2_6/ptodata/l/ina/6A_COMB.seq:*
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US-09-245-041-5
US-09-342-681C-8
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US-09-097-319A-19
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US-08-799-113-9
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US-09-801-052-3
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US-09-221-017B-372
US-09-235-564-4
US-08-241-387-4
US-08-241-387-2
US-09-627-769-1
US-09-627-769-1
US-09-627-769-1
US-09-627-769-1
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570.891 Million cell updates/sec
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                                            e 19, Apple e 9, Appli e 9, Appli e 11, Appli e 11, Appli e 11, Appli e 20, Appli e 21, Appli e 21, Appli e 21, Appli e 21, Appli e 3, Appli e 
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e 5, Appli
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RESULT 1		c 45	C 44	c 43	c 42	c 41			c 38			c 35	34		c 32	31	c 30	c 29	28
		24.6	24.6	24.6	24.6	24.6	24.6	24.6	24.6	24.8	24.8	24.8	24.8	25	25	25	25	25	25
		20.3	20.3	20.3	20.3	20.3	20.3	20.3	20.3	20.5	20.5	20.5	20.5	20.7	20.7	20.7	20.7	20.7	20.7
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	ALIGNMENTS	US-08-328-809-3	US-08-258-442-8	US-07-814-964-8	PCT-US92-11107-9	US-08-328-809-4	US-08-258-442-9	US-07-814-964-9	US-08-457-752-2	US-08-458-434A-6	US-08-646-715-10	US-08-188-582-10	US-08-347-471-1	US-08-705-557-1	US-09-426-436-1	US-09-428-517-1	US-09-075-904-1	US-08-247-901C-1	US-08-961-527-124
		Sequence 3, Appli	Sequence 8, Appli	Sequence 8, Appli	Sequence 9, Appli	Sequence 4, Appli	Sequence 9, Appli	Sequence 9, Appli	Sequence 2, Appli	Sequence 6, Appli	Sequence 10, Appl	Sequence 10, Appl	Sequence 1, Appli	Sequence 124, App					

US-09-750-580-4

Patent No. 6455280
GENERAL INFORMATION:

Sequence 4, Application US/09750580 Patent No. 6455280

APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:

Bihain, Bour, Barbara Denison, Blake Frances

Bernard

Dumas Milne Edwards, Jean-Baptiste

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APPLICANT: Duclert, Aymeric
APPLICANT: Bouqueleret, Lydie
APPLICANT: Bouqueleret, Lydie
APPLICANT: Bouqueleret, Lydie
APPLICANT: Salter-Cid, Luisa
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITING NEOPLASTIC CELL GROWT
FILE REFERENCE: 89.USZ.CIP
CURRENT APPLICATION NUMBER: US/09/750,580
CURRENT APPLICATION NUMBER: US 09/599,362
PRIOR APPLICATION NUMBER: US 09/599,362
PRIOR FILING DATE: 2000-06-21
PRIOR FILING DATE: 2000-06-21
PRIOR APPLICATION NUMBER: PCT/IB00/0101
PRIOR APPLICATION NUMBER: PCT/IB09/0258
PRIOR APPLICATION NUMBER: US 49/469/099
PRIOR FILING DATE: 1999-12-20
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: US 60/141,032
PRIOR APPLICATION NUMBER: US 60/141,032
PRIOR FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Pate SEQ ID NO 4 LENGTH: 5381
                    NAME/KEY: exon
LOCATION: 1613..1724
OTHER INFORMATION: exon
                                                                                                          LOCATION: 1442..1498
OTHER INFORMATION: exon
                                                                                                                                                                                                 LOCATION: 919..930
OTHER INFORMATION: exon
                                                                                                                                                                                                                                                                           LOCATION: 1..918
OTHER INFORMATION: 5'regulatory region
                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                      NAME/KEY: exon
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NAME/KEY:
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LOCATION: 3941..5381
OTHER INFORMATION: 3'regulatory region
                                                                                                                                                                                                                                                                                                       NAME/KEY: primer_bind
LOCATION: 3214..3232
OTHER INFORMATION: 17-41-250.mis complement
NAME/KEY: misc_binding
LOCATION: 307..331
OTHER INFORMATION: 17-42-319.probe
NAME/KEY: misc_binding
LOCATION: 3201..3225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: primer_bind
LOCATION: 553..11575
OTHER INFORMATION: 17-42.rp complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: conflict
LOCATION: 1241
OTHER INFORMATION: 17-39-343 : T in ref genbank AC007707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: allele
LOCATION: 3213
OTHER INFORMATION: 17-41-250 : polymorphic base C or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: allele
LOCATION: 319
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OTHER INFORMATION: exon 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 300..318
OTHER INFORMATION: 17-42-319.mis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 3432..14454
OTHER INFORMATION: 17-41.rp complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: primer_bind LOCATION: 2964..13984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: 17-40.rp complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: 17-39.rp complement
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OTHER INFORMATION: 17-40.pu
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LOCATION: 1246..12267
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OTHER INFORMATION: 17-39.pu
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LOCATION: 899..11920
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LOCATION: 1..110:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: 17-42-319 : polymorphic base C or T
                                                                                                                                                                                                                                                                                       OTHER INFORMATION: 17-41-250.probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: 17-41-250.mis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: 17-42-319.mis complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: primer_bind LOCATION: 300..318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: primer_bind LOCATION: 3432..14454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: 17-41.pu
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LOCATION: 1632..12651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: primer_bind
LOCATION: 1441..12461
312 CAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCTTGC
                                                                                        252
                         61 CAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCTTGC 120
                                                                                                                                                                                                   Local Similarity
                                                                                                               1 GCATTTGGGCTTTGCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGG 60
                                                                                      GCATTTGGGCTTGCTCCTCAGAGGCCCCTGCGAGTTGGAGTTCAGCTTTTCCTCATGGGG 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer_bind
3194..3212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer_bind
                                                                                                                                                                       99.7%; Score 120.6; DB ilarity 100.0%; Pred. No. 2e-30; Conservative 0; Mismatches
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                                                                                                                                                                                                                   Length 5381;
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US-09-750-580-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
SEQ ID NO 1
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PRIOR FILING DATE: 1999-1
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1998-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/750,580
CURRENT FILING DATE: 2000-12-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bougueleret, Lydie
APPLICANT: Ebbets-Reed, Dana
APPLICANT: Salter-Cid, Luisa
APPLICANT: Salter-Cid, Luisa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2000-PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
NAME/KEY: misc_feature
LOCATION: 10946..12946
OTHER INFORMATION: 5'regulatory region
NAME/KEY: exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapiens
            NAME/KEY: allele LOCATION: 42218 OTHER INFORMATION:
                                                           NAME/KEY: allele
LOCATION: 15241
OTHER INFORMATION:
                                                                                                                                                                                                                           OTHER INFORMATION: exon NAME/KEY: misc_feature LOCATION: 15969..17969
                                                                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
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OTHER INFORMATION: exon
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OTHER INFORMATION: exon
                                                                                                              OTHER INFORMATION:
                                                                                                                              NAME/KEY: allele LOCATION: 12347
                                                                                                                                                               OTHER INFORMATION:
                                                                                                                                                                             NAME/KEY: allele
LOCATION: 1239
                                                                                                                                                                                                              OTHER INFORMATION: 3'regulatory region
                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: exon
                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: exon
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                                                                                                                                                                                                                                                                                 14271..15968
                                                                                                                                                                                                                                                                                                                                13641..13752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dumas Milne Edwards,
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Bour, Barbara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bihain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Duclert, Aymeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yen, Frances
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ON: METHODS AND COMPOSITIONS FOR INHIBITING NEOPLASTIC CELL, GROWT 89.US2.CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1999-12-21
               20-841-149
                                                                                                                                                            20-828-311 : polymorphic base C or T
                                                             17-41-250
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              : polymorphic base A or
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                                                             polymorphic base C or
                                                                                                              polymorphic base C or
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NAME/KEY: primer_bind
LOCATION: 42572..42591
OTHER INFORMATION: 20-841.rp
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LOCATION: 12581..12603
OTHER INFORMATION: 17-42.rp complement
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LOCATION: 77166..77185
OTHER INFORMATION: 20-853.rp
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LOCATION: 45328..45347
OTHER INFORMATION: 20-842.pu
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                                                                                                                                                        OTHER INFORMATION: 20-841-149.mis NAME/KEY: primer_bind LOCATION: 42219.42237
                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: primer_bind
LOCATION: 12348..12366
OTHER INFORMATION: 17-42-319.mis complement
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LOCATION: 1240..1258
OTHER INFORMATION: 20-828-311.mis complement
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LOCATION: 15242..15260
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OTHER INFORMATION: 17-42-319.mis
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OTHER INFORMATION: 20-828-311.mis
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OTHER INFORMATION: 20-842-115.mis complement
                       NAME/KEY: primer_bind LOCATION: 45443..45461
                                                                  OTHER INFORMATION: 20-842-115.mis
                                                                                        NAME/KEY: primer_bind LOCATION: 45423..45441
                                                                                                                                   OTHER INFORMATION: 20-841-149.mis complement
                                                                                                                                                                                                                           NAME/KEY: primer_bind
LOCATION: 42199..42217
                                                                                                                                                                                                                                                                      OTHER INFORMATION: 17-41-250.mis complement
                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 17-41-250.mis
                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: primer_bind LOCATION: 15222..15240
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LOCATION: 76644..76664
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LOCATION: 45863..45883
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12328..12346
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                                                                                                                                                     ; TYPE: DNA; ORGANISM: Mus musculus US-09-245-041-5
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                                                                                                                                                                                                            SOFTWARE: Fast:
SEQ ID NO 5
LENGTH: 90050
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Best Local Similarity
                                                                                             Query Match
Best Local
                                                                            Matches
                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/245,041
CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/093,630
EARLIER FILING DATE: 1998-07-21
EARLIER APPLICATION NUMBER: 60/104,978
EARLIER FILING DATE: 1998-07
EARLIER FILING DATE: 1998-07
NUMBER OF SEQ ID NOS: 131
                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND TREATMENT TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY FILE REFERENCE: 7853-136
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Moore, K. APPLICANT: Nagle, D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: 20-853-415.mis

NAME/KEY: primer_bind
LOCATION: 77059..77077

OTHER INFORMATION: 20-853-415.mis complement
NAME/KEY: misc_binding
LOCATION: 1227..1251
12400 C 12400
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LOCATION: 77046..77070
OTHER INFORMATION: 20-853-415.probe
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                                                                                             Similarity
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57.8%;
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100.0%;
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                                                                                             Score 27; DB 4;
Pred. No. 25;
                                                                            Mismatches
                                                                            35;
                                                                                                               Length 90050;
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32998 GTCCCAGGGGATCTGATGCCCAG 32976

81 GTTCCTGGGGCTCAGAGTCCCTG 103

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 6384207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/342,681C CURRENT FILING DATE: 1999-06-29 PRIOR ENTITY OF THE PRIOR APPLICATION NUMBER: 60/92,279 PRIOR FILING DATE: 1998-07-09 PRIOR APPLICATION NUMBER: 60/112,366 PRIOR FILING DATE: 1998-12-15 PRIOR FILING DATE: 1998-12-15 PRIOR FILING DATE: 1998-12-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Zonana et al. TITLE OF INVENTION: Hypohydrotic ectodermal dysplasia genes and proteins FILE REFERENCE: 52978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Ver.
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 977
                                 COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/097,319A
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                    APPLICANT: Woosley, Aaron
TITLE OF INVENTION: Regulatory Sequences for Transgenic Plants
NUMBER OF SEQUENCES: 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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CITY: Indianapolis
STATE: Indiana
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CLASSIFICATION: 800
                    FILING DATE:
                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                              46268
                                                                                                                                                                                                                                                         9330 Zionsville Road
                                                                                                                                                                                                 USA
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Petolino, Joseph F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Belmar, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Armstrong, Katherine
                                                                                                                                                                                                                                                                                                                                                                             Smith, Kelley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ainley, Michael
                                                                                                                                                                                                                                                                            DowElanco Patent Department
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    Mismatches

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; NUMBER OF SEQ ID NOS: 54
; SOFTWARE: Microsoft Word Version 7.0A
; SEQ ID NO 52
; LENGTH: 995
; TYPE: DNA
; ORGANISM: Triticum aestivum
US-09-173-300-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
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US-09-173-300-52/c
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                                                                                                                                                                                                                                                                        RESULT 7
US-08-583-562B-9
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                                                                                                                                                                                                                      Sequence 9, Application US/08583562B Patent No. 5922570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT: Hitz, William D.
APPLICANT: Kinney, Anthony J.
APPLICANT: Kinney, Anthony J.
APPLICANT: Rafalski, J. Antoni
TITLE OF INVENTION: PLANT BRANCHED CHAIN AMINO ACID BIOSYNTHETIC
FILE REFERENCE: BB-1126
FILE REFERENCE: BB-1126
CURRENT FILING DATE: 1998-10-15
EARLIER APPLICATION NUMBER: 60/063,423
EARLIER APPLICATION NUMBER: 60/063,423
EARLIER FILING DATE: 1997 October 28
NUMBER: 650 15 NOC: 54

EARLIER FILING DATE: 1997 October 28
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             GENERAL INFORMATION:

APPLICANT: Staunton, Donald

APPLICANT: Staunton, Donald

APPLICANT: Harris, Edith

TITLE OF INVENTION: Cytoplasmic Modulators of Integrin

TITLE OF INVENTION: Binding

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:

ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun

ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
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TELEPHONE: 317 337 4816
TELEFAX: 317 337 4847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: C: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5920 GGCATGTCCGCCTCGGTGGCACGGCGGATGTCTGGGGGCCATGGCT 5861
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LENGTH: 9335 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION: NAME: Stuart, Donald R
                                                                                                                                                                                                                                                                                                                                                                          778 CTTCATCTCCGCAGAGGCTCGGCTGCATAAGCTGATCTTTCCCTCATGCAGCAGCTTTC 720
                                                                                                                                                                                                                                                                                                                                                                                                                          10 CTTGCTCTCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTY 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y Match 22.0%; Score 26.6;
Local Similarity 64.4%; Pred. No. 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68 YACTTTCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCT 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: double
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  STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               38;
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233 South Wacker Drive, 6300 Sears Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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COMPUTER READABLE FORM:

CITY: Chicago STATE: Illino:

Illinois

COUNTRY:

USA

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SEQUENCE CHARACTERISTICS:
LENGTH: 1125 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Williams Jr., Joseph A. REGISTRATION NUMBER: 38.659 REFERENCE/DOCKET NUMBER: 2786 TELECOMMUNICATION INFORMATION: TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, VICURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/583,562B
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Staunton, Donald E. APPLICANT: Harris, Edith S. TITLE OF INVENTION: Cytoplasmic Modulators of Integrin TITLE OF INVENTION: Binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Williams Jr., Jose
                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 65 CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
NAME: Greta E. No. 5948891and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        414 CCGGTTTGTG 423
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CTTY: Chicago
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COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Local Similarity 60.0%;
nes 42; Conservative
                                          CLASSIFICATION: 536
                                                                                                                                                                                                                                                         COUNTRY:
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                                                                  FILING DATE:
                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                       60606
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                                                                                                                                                                                                                                                         United States of America
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                                                                                                                                                                                                                                                                                                                                             Marshall, O'Toole, Gerstein, Murray & Borun
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                                                                                      us/08/779,113

    Mismatches

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US-08-779-113-9
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INFORMATION FOR SEQ ID NO: 9:
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                                                                                                                                                   INFORMATION FOR SEQ ID NO:
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REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Staunton, Donald APPLICANT: Harris, Edith TITLE OF INVENTION: Cytoplas TITLE OF INVENTION: Binding
                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Williams Jr. JOSEPH A.
REGISTRATION NUMBER: 38,659
REFERENCE/DOCKET NUMBER: 2786
                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 2-STREET: Chicago
CITY: Chicago
TTATE: Illinois
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                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 1494 base pairs
                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                 FEATURE:
                                                 MOLECULE TYPE:
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                                                                                                                                                               TELEFAX:
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                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                             COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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COMPUTER: 1F
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NAME/KEY:
LOCATION:
                                                                TOPOLOGY:
                                                                             STRANDEDNESS:
                                                                                                  TYPE:
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233 South Wacker Drive, 6300 Sears Tower
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                                                                linear
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US-08-779-113-11
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                                                                                                                                                Query Match
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Best Local Similarity
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APPLICANT: Staunton, Donald
APPLICANT: Harris, Edith S.
TITLE OF INVENTION: Cytoplas
TITLE OF INVENTION: Binding
                                                                                                                                                                                                                                                                                                                                 TELEFAX: 312-474-0448
INFORMATION FOR SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 312-474-6300
                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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                                                     78 CCAGTTCCTG 87
                                                                                                                                  Local Similarity
                                                                                                                                                                                                             NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                              NAME: Greta E. No. 5948891and REGISTRATION NUMBER: 35,302 REFERENCE/DOCKET NUMBER: 2786
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                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/779,113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
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                                                                                                                                                                                                                                                                                                  ENGTH:
                                                                                                                   42; Conservative
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                                                                                                                                                                                                                                                                                                  1494 base pairs
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233 South Wacker Drive, 6300 Sears Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             United States of America
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                                                                                                                              21.8%;
60.0%;
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    Mismatches

                                                                                                                                  Score 26.4; DB 2; Pred. No. 11;
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Pred. No. 11;
                                                                                                                  27; Indels
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US-08-779-113-1
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US-08-583-562B-1
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                                                                                                                                                                                                          Sequence 1, Application US/08779113 Patent No. 5948891
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                                                                GENERAL INFORMATION:
APPLICANT: Staunton, Donald E.
APPLICANT: Harris, Edith S.
TITLE OF INVENTION: Cytoplasmic Modulators of Integrin
TITLE OF INVENTION: Binding
NUMBER OF SEQUENCES: 65
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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TELEFAX: 312-474-0448
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                            1230 CCGGTTTGTG 1239
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LENGTH: 2574 base pairs
TYPE: nucleic acid
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NAME: Williams Jr., Joseph A.
REGISTRATION NUMBER: 38,659
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                  1170 CCCCTCTGGCCGTGTGACTGAGGCAGAGATTGTGCCCATGGGGAAGAACTCACACTGCGT 1229
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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STREET: 200
STREET: 200
Chicago
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                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/583,562B
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STATE: Illinois
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                                    E: Marshall, O'Toole, Gerstein, Murray & Borun 233 South Wacker Drive, 6300 Sears Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Marshall, O'Toole, Gerstein, Murray & Borun
233 South Wacker Drive, 6300 Sears Tower
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    Mismatches

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Pred. No. 13;
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COMPUTER READABLE FORM:

COMPUTER:

MEDIUM TYPE:

Floppy disk

COUNTRY: United States of America ZIP: 60606

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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/779,113
                                  ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Megabase Transcript Map: No. 58: TITLE OF INVENTION: Sequences and Antibodies Thereto NUMBER OF SEQUENCES: 31
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NAME: Greta E. No. 5948891and
REGISTRATION NUMBER: 35,302
                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Tsuchihashi, Zenta APPLICANT: Wolff, Roger K.
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APPLICATION NUMBER: US/01 FILING DATE: 01-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            42;
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                    US/08/724,394A
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Db 160999 GCCGTTCGCCTATTTCTTGGCCGCCCTCTTAGGCTTGACAACCTTGGCCTTAGC 161058
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US-08-724-394A-21
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INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
SEQUENCE C46240 base pairs
INFORMATION FOR SEQ ID NO:
                                  ATTORNEY/AGENT INFORMATION:
NAME: FILLS, Renee A.
REGISTRATION NUMBER: 35,136
REFERENCE/DOCKET NUMBER: 011
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/724,394A
FILING DATE: 01-OCT-1996
CLASSIFICATION: 536
CLASSIFICATION: 536
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TELEPHONE: 415-576-0200
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                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: TOWNSEND and TOWNSEND and CREW LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Wolff, Roger K.
TITLE OF INVENTION: Megabase Transcript Map: No. 58:
TITLE OF INVENTION: Sequences and Antibodies Thereto
NUMBER OF SEQUENCES: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37 GGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTCCTGGGGCTCAGA 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: not r
TOPOLOGY: not relev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97 GTCCCTGGCC 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature LOCATION: 1..246240 OTHER INFORMATION: /not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 35
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                       San Francisco
                                                                                                                                                                                                                                                                                                                                                                                    CA
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                                                                                                                                                                                                                                                                                                                                                               USA
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Ruddy, David A.
                   415-576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Feder, John N.
Kronmal, Gregory S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thomas, Winston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                            017957-000100
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RESULT 15
US-08-724-394A-22
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; OTHER INFORMATION: /note= "HLA-H.CONTIG"
US-08-724-394A-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Db 161059 GGCCTTGGGC 161068
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22, Application US/08724394A Patent No. 5872237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Feder, John N.
APPLICANT: Kronmal, Gregory S.
APPLICANT: Lauer, Peter M.
                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
TELEFAX: 415-576-0300
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 246240 base pairs
                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Rclease #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/724,394A FILING DATE: 01-0CT-1996
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-100S
                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Fitts, Renee A.
REGISTRATION NUMBER: 35,136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Megabase Transcript Map: No. 5872237el TITLE OF INVENTION: Sequences and Antibodies Thereto NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ruddy, David A.
APPLICANT: Thomas, Winston
APPLICANT: Tsuchhashi, Zenta
APPLICANT: Wolff, Roger K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 246240 base pa
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                                            MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Two Embarca
CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
  NAME/KEY:
                                                                                 STRANDEDNESS:
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                                                                   TOPOLOGY:
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                                                                                                        nucleic acid
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Ruddy, David A.
                                                              ss: not relevant not relevant
misc_feature
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Title:
Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
                                                          \circ \circ \circ \circ \circ \circ \circ
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                                                                                                                                                                                                                                                                                                                                                                  Score
        Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-842-364-1_COPY_12280_12400
121
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        : /cgn2_6/ptodata//pubpna/US07_PUBCOMB.seq:*
:/cgn2_6/ptodata//pubpna/US07_PUBCOMB.seq:*
:/cgn2_6/ptodata//pubpna/US06_PUBCOMB.seq:*
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:/cgn2_6/ptodata//pubpna/US07_NEW_PUB.seq:*
:/cgn2_6/ptodata//pubpna/US08_NEW_PUB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                  Length DB
                                                                                               317
504
1026
1091
3000
222
524
4926
90050
1147
15980
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    10 US-09-751-877-4
11 US-09-751-877-1
11 US-09-718-446-405
11 US-09-718-45-405
11 US-09-718-50-553
11 US-09-816-055-28
12 US-09-919-177-13
12 US-09-919-177-13
13 US-09-738-550-11626
14 US-09-788-95-18
15 US-09-788-95-18
16 US-09-788-95-18
17 US-09-784-864-7714
18 US-09-917-800A-61
19 US-09-917-800A-61
19 US-09-917-800A-61
10 US-09-917-800A-61
10 US-09-917-800A-61
10 US-09-917-805-52
10 US-09-9184-761-25624
10 US-09-864-761-25624
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Sequence 4, Appli
Sequence 1, Appli
Sequence 5953, App
Sequence 28, Appli
Sequence 1, Appli
Sequence 11626, A
Sequence 11824, A
Sequence 15294, A
Sequence 15, Appli
Sequence 5, Appli
Sequence 5, Appli
Sequence 5116, Appli
Sequence 5116, Appli
Sequence 5116, Appli
Sequence 52, Appli
Sequence 53, Appli
Sequence 52, Appli
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Sequence 53, Appli
Sequence 25624, Appli
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	24	24.8	24.8	24.8	25	25	25.2	25.2	25.2	25.2	25.2	25.2	25.2	25.2	25.2	25.4	25.6	25.6	26	26	26	26	26	26.4	26.4	26.4
	ر د د	20.5	20.5	20.5	20.7	20.7	20.8	20.8	20.8	20.8	20.8	20.8	20.8	20.8	20.8	21.0	21.2	21.2	21.5	21.5	21.5	21.5	21.5	21.8	21.8	21.8
	1837	1827	1632	261	720	720	5762	5760	5629	5265	5265	674	674	424	350	4744	3741	470	1416	1206	1157	1151	1138	5607	1391	1391
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	115-09-919-172-48	US-09-974-298-174	US-09-887-38: 3	US-09-728-444-1027	US-09-815-242-9471	US-09-815-242-9319	US-09-919-172-55	US-10-044-090-67	US-09-969-347-208	US-10-033-528-1663	US-09-920-300A-1663	US-09-764-860-184	US-09-764-878-47	US-09-867-550-705	US-09-878-574-3008	US-09-880-107-3459	US-09-803-286A-1	US-09-864-761-11312	US-10-062-254-145	US-10-062-254-143	US-09-778-844-166	US-09-778-844-164	US-09-778-844-165	US-10-108-605-92	US-09-880-107-3740	US-09-969-708-480
		Sequence 174, App	Sequence 3, Appli	Sequence 1027, Ap	Sequence 9471, Ap	Sequence 9319, Ap	Sequence 55, Appl	Sequence 67, Appl	Sequence 208, App	Sequence 1663, Ap	Sequence 1663, Ap	Sequence 184, App	Sequence 47, Appl	Sequence 705, App	Sequence 3008, Ap	Sequence 3459, Ap	Sequence 1, Appli	Sequence 11312, A	Sequence 145, App	Sequence 143, App	Sequence 166, App	Sequence 164, App	Sequence 165, App	Sequence 92, Appl	Ó	Sequence 480, App

## ALIGNMENTS

RESULT 1 US-09-751-877-4

Sequence 4, Application US/09751877 Patent No. US20020142949A1

INFORMATION:

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GENERAL INFORMATION:

APPLICANT: Pen, Frances
APPLICANT: Denison, Blake
APPLICANT: Denison, Blake
APPLICANT: Biháin, Bernard
APPLICANT: Biháin, Bernard
APPLICANT: Bour, Barbara
APPLICANT: Dunds Milne Edwards, Jean-Baptiste
APPLICANT: Dunds Milne Edwards
APPLICANT: Bougueleret, Lydie
APPLICANT: Bougueleret, Lydie
APPLICANT: Bougueleret, Lydie
APPLICANT: Salter-Cid, Luisa
APPLICANT: Salter-Cid, Luisa
APPLICANT: Salter-Cid, Luisa
APPLICANT: Salter-Cid, Luisa
CURRENT: APPLICATION MEMBER: US/09/751,877
CURRENT FILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patent.pm
SEQ ID NO 4
LENGTH: 5381
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/REY: misc_feature
LOCATION: 1.918
OTHER INFORMATION: 5'regulatory region
NAME/REY: exon
LOCATION: 919.930
OTHER REPERANTON: Exon 1
                 NAME/KEY: exon
LOCATION: 1442..1498
OTHER INFORMATION: exon 2
NAME/KEY: exon
LOCATION: 1613..1724
OTHER INFORMATION: exon 3
NAME/KEY: exon
LOCATION: 2243..3940
OTHER INFORMATION: exon 4
                                                                             w
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: NAME/KEY: primer_bind
: IOCATION: 3214..3232
: OTHER INFORMATION: 17-41-250.mis complement
: NAME/KEY: misc_binding
: LOCATION: 307..331
: OTHER INFORMATION: 17-42-319.probe
: NAME/KEY: misc_binding
: LOCATION: 3201..3225
: OTHER INFORMATION: 17-41-250.probe
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OTHER INFORMATION: 17-39-343 : T in ref genbank aC007707
NAME/KEY: CONFLICT
LOCATION: 1447
OTHER INFORMATION: 17-40-202 : G in ref genbank aC007707
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LOCATION: 3432..14454
OTHER INFORMATION: 17-41.rp complement
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LOCATION: 1632..12651
OTHER INFORMATION: 17-40.rp complement
NAME/KEY: primer_bind
LOCATION: 2564.13984
OTHER INFORMATION: 17-41.pu
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LOCATION: 1441..12461
OTHER INFORMATION: 17-39.rp complement
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LOCATION: 899..11920
OTHER INFORMATION: 17-39.pu
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LOCATION: 553..11575
OTHER INFORMATION: 17-42.rp complement
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LOCATION: 1..11022
OTHER INFORMATION: 17-42.pu
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LOCATION: 319

OTHER INFORMATION: 17-42-319: polymorphic base C or T
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LOCATION: 320..338
OTHER INFORMATION: 17-42-319.mis complement
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LOCATION: 300..318
OTHER INFORMATION: 17-42-319.mis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 17-41-250 : polymorphic base C or T
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LOCATION: 3213
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LOCATION: 3194..3212
OTHER INFORMATION: 17-41-250.mis
                                                                     372 C 372
                                  121 C 121
                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                          Similarity 100.
21; Conservative
                                                                                                                                                                                                                                         99.7%; Score 120.6; DB 10; Length 5381; 100.0%; Pred. No. 9.3e-31;

 Mismatches

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                                                                                                                                                                                                                     Indels
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RESULT 2
US-09-751-877-1
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APPLICANT: Yen, Fra-
APPLICANT: Denison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: 10946..12946
OTHER INFORMATION: 5'regulatory region
NAME/KEY: exon
LOCATION: 12947. 12958
OTHER INFORMATION: exon 1
NAME/KEY: exon
LOCATION: 13470..13526
OTHER INFORMATION: exon 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Bour, Barbara
APPLICANT: Bihain, Bernard
APPLICANT: Dumas Milne Edwards, Jean-Baptiste
APPLICANT: Dumas Milne Edwards, Jean-Baptiste
APPLICANT: Bougueleret, Lydie
APPLICANT: Bobets Reed, Dana
APPLICANT: Salter-Cid, Luisa
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITING NEOPLASTIC CELL GROWTH
FILE REFERENCE: 89.US3.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09751877 Patent No. US20020142949A1
                                                                                                                                                                                                                                                                                 NAME/KEY: allele
LOCATION: 15241
OTHER INFORMATION: 17-41-250 :
NAME/KEY: allele
LOCATION: 42218
OTHER INFORMATION: 20-841-149 :
NAME/KEY: allele
LOCATION: 45442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/751,877
CURRENT FILING DATE: 2000-12-28
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patent.pm
EQ ID NO 1
LENGTH: 81001
NAME/KEY: primer_bind
LOCATION: 12029..12050
OTHER INFORMATION: 17-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: allele
LOCATION: 1239
COTHER INFORMATION: 20-828-311 : polymorphic base C
NAME/KEY: allele
                                                             NAME/KEY: primer_bind
LOCATION: 1357..1377
OTHER INFORMATION: 20-828.rp
                                                                                                                             NAME/KEY: primer_bind LOCATION: 929.949 OTHER INFORMATION: 20-828.pu
                                                                                                                                                                                               OTHER INFORMATION: 20-842-115
NAME/KEY: allele
LOCATION: 77058
OTHER INFORMATION: 20-853-415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 12347
OTHER INFORMATION: 17-42-319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: 15969..17969
OTHER INFORMATION: 3'regulatory region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: exon
LOCATION: 14271..15968
OTHER INFORMATION: exon 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: exon
LOCATION: 13641..13752
OTHER INFORMATION: exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Denison, Blake
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Frances
  17-42.pu
                                                                  complement
                                                                                                                                                                                                                                                                  : polymorphic base A or
                                                                                                                                                                                                                                                                                                                                  : polymorphic base A or
                                                                                                                                                                                                  : polymorphic base C or
                                                                                                                                                                                                                                                                                                                                                                                                  : polymorphic base C or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : polymorphic base C or T
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NAME/KEY: misc_binding
LOCATION: 1235.12359
OTHER INFORMATION: 17-42-319.probe
NAME/KEY: misc_binding
LOCATION: 15229.15233
OTHER INFORMATION: 17-41-250.probe
NAME/KEY: misc_binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: primer_bind
LOCATION: 42070..42090
OTHER INFORMATION: 20-841.pu
NAME/KEY: primer_bind
LOCATION: 42572.42591
OTHER INFORMATION: 20-841.rp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: primer_bind LOCATION: 45863..45883 OTHER INFORMATION: 20-84 NAME/KEY: primer_bind LOCATION: 76644..76664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: primer_bind
LOCATION: 1240..1258
OTHER INFORMATION: 20-828-311 min
                                                                                                                                                                                                                                                    NAME/KEY: primer_bind
LOCATION: 77039. 77057
OTHER INFORMATION: 20-853-415.mis
NAME/KEY: primer_bind
LOCATION: 77075. 77077
OTHER INFORMATION: 20-853-415.mis complement
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LOCATION: 15460..15482
OTHER INFORMATION: 17-41.rp complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: primer_bind
LOCATION: 15222..15240
OTHER INFORMATION: 17-41-250.mis
NAME/KEY: primer_bind
LOCATION: 15242..15260
OTHER INFORMATION: 17-41-250.mis complement
NAME/KEY: primer_bind
LOCATION: 42199.42217
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LOCATION: 45328..45347
OTHER INFORMATION: 20-842.pu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: 17-42.rp complement NAME/KEY: primer_bind LOCATION: 14992..15012
                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: primer_bind
LOCATION: 45423..45441
OTHER INFORMATION: 20-
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LOCATION: 42219. 42237
OTHER INFORMATION: 20-841-149.mis complement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: primer_bind
LOCATION: 12328..12346
OTHER INFORMATION: 17-42-319.mis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 20-6
NAME/KEY: primer_bind
LOCATION: 77166..77185
OTHER INFORMATION: 20-6
                                                                                                                                                                                              OTHER INFORMATION:
                                                                                                                                                                                                               NAME/KEY: misc_binding LOCATION: 1227..1251
                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: primer_bind LOCATION: 45443..45461
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION:
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LOCATION: 12348..12366
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION:
20-841-149.probe
                                                                                                                                                                                         20-828-311.probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-841-149.mis
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RESULT 4

US-09-783-590-5953
Sequence 5953, Application US/09783590
Patent No. US20020110850A1
GENERAL IMPORMATION:
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:

APPLICANT: Friedrich, Glenn
APPLICANT: Friedrich, Glenn
APPLICANT: Sands Arthur T.

ITILE OF INVENTION: No. US2020081668Alel Murine Polynucleotide Sequences
ITILE OF INVENTION: and Mutant Cells and Mutant Animals Defined Thereby
FILE REFERENCE: LEX-0101-USA
CURRENT APPLICATION NUMBER: US,709/728,446
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/168,270
PRIOR FILING DATE: 1999-12-01
UNDBER OF SEQ ID NOS: 1461
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 405
LENGTH: 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
US-09-728-446-405
US-09-728-446-405
PEROPERTY 3
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LOCATION: 45430...4544
OTHER INFORMATION: 20-842-115.probe
NAME/KEY: misc_binding
LOCATION: 77046..77070
OTHER INFORMATION: 20-853-415.probe
US-09-751-877-1
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; ORGANISM: Mus
US-09-728-446-405
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Best Local S
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Best Local Similarity
Matches 121; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12400 C 12400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121 C 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80 AGTTCCTGGGGCTCAGAGTCCCTGGCCCAG 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 Match 8 Matches 7 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 TCAGAGGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCC 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGTTCCAGAGGAATTGACACCCTTGGACAG 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09728446
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR FILING DATE: 1995-04-12
PRIOR FILING DATE: 1994-11-21
PRIOR FILING DATE: 1994-11-21
UMBER OF SEO ID NOS: 12485
SOFTWARE: PALENTIN VET. 2.0
SEQ ID NO 5933
LENGTH: 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COCATION: (61)

OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (62)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (78)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (96)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (115)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (115)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (126)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (126)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (126)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (127)
LOCATION: (126)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (127)
LOCATION: (127)
LOCATION: (127)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (127)
LOCATION: (127)
                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (153)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (164)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (170)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc feature
LOCATION: (130)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (132)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (146)
                NAME/KEY: misc feature
LOCATION: (240)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (257)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (146)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (150)
LOCATION: (150)
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ORGANISM: Homo sapiens
                                                                                                                                                        LOCATION: (188)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (199)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COCATION: (150)
OTHER INFORMATION: n equals a,t,g, or c
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc feature
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; OTHER INFORMATION: n equals a,t,g, or c US-09-783-590-5953
                                                                                                                                                                                                                                                                  COCATION: (461)

OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (462)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (470)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (486)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (486)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (496)
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NAME/KEY: misc feature
LOCATION: (427)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (429)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LOCATION: (439)
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LCCATION: (443)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LCCATION: (456)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
LCCATION: (457)
OTHER INFORMATION: n equals a,t,9, or c
NAME/KEY: misc feature
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LOCATION: (303)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (364)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (399)
          Matches
                                                          Query Match
Best Local
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LOCATION: (406)
OTHER INFORMATION: n equals a,t,g, or
NAME/KEY: misc feature
LOCATION: (421)
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LOCATION: (288)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (439)
OTHER INFORMATION: n equals a,t,9,
NAME/KEY: misc feature
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OTHER INFORMATION: n equals a,t,9, or
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     Local Similarity es 44; Conserv
23.0%;
ilarity 57.1%;
Conservative
; Score 27.8; DB 10;
; Pred. No. 2.9;
1; Mismatches 32;
     Indels
                                                                                         Length 504;
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225 TTACTGGGAGGCAGTNT 241

82 TTCCTGGGGCTCAGAGT

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US-09-886-055-28; Sequence 28; A
                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-011-597-1
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                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/011,597
CURRENT FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US/09/353,243
PRIOR FILING DATE: 1999-07-14
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.0 - beta
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: BORINI, James A.
APPLICANT: BORINIS JAMES A.
APPLICANT: BORONSKY, Beth E.
TITLE OF INVENTION: DNA ENCODING ORPHAN SNORF40 RECEPTOR
FILE REFERENCE: 59587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/886,055
CURRENT FILLING DATE: 2001-06-22
PRIOR APPLICATION UNMBER: 60/213,812
PRIOR FILLING DATE: 2000-06-22
NUMBER OF SEQ 1D NOS: 522
SOFTWARE: Patentin Ver. 2.1
SEQ 1D NO 28
SEQ 1D NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/10011597 Patent No. US20020161217A1
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APPLICANT: ZOZULYA, SERGEY
TITLE OF INVENTION: RECEPTOR FINGERPRINTING, SENSORY PERCEPTION, AND
TITLE OF INVENTION: BIOSENSONS OF CHEMICAL SENSANTS
FILE REFERENCE: 078003-0277150
                                                                                                                                                                                                                     Query Match
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123 GATGAACTCAGAGAACCTCACCCGGGCCGCGGTTGCC 159
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                                           85 CTGGGGCTCAGAGTCCCTGGCCCAGATGCCTCTTGCC 121
                                                                                      63 GCCTTGCCATTGCTCTTATCTCCCTCCTGCTTTGCCTCTCTCAGTCTCTGTCCAGTAG 122
                                                                                                                              25 GGCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYACTTTCGCTCCAGTTC 84
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                                                                                                                                                                                              Similarity 54.6%;
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Similarity 54.6%;
                                                                                                                                                                              Conservative
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                                                                                                                                                                         Score 27.8; DB 9; Length 1091; Pred. No. 3.8; 1; Mismatches 43; Indels 0

    Mismatches

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GENERAL INFORMATION:

APPLICANT: Faris, Mary
APPLICANT: Faris, Mary
APPLICANT: Turner, Christopher M.

TITLE OF INVENTION: PROSTATE CANCER MARKERS
FILE REFERENCE: PA-0036 US
CURRENT APPLICATION NUMBER: US/09/919,172
CURRENT FILL NG DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/222,469
PRIOR FILLING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 102
SOFTMARE: PERL PROGRAM
SEQ ID NO 13
LENGTH: 3000
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US-09-919-172-13/c
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; OTHER INFORMATION: Incyte ID No. US20020119463A1 995610.1
US-09-919-172-13
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APPLICANT: Ruben, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT APPLICATION NUMBER: 08/420,856
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
RUMBER OF SEQ ID NOS: 12465
SOFTMARE: Patentin Ver: 2.0
SEQ ID NO 11626
LENGTH: 222
ORGANISM: Homo sapiens
FEARTURE:
NAME/KEY: misc feature
LOCATION: (1)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (5)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (12)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (25)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (25)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
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Patent No. US20020110850A1
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Best Local S
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APPLICANT: Haseltine, William A.
APPLICANT: Li, Haddong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR TILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00661
PRIOR APPLICATION NUMBER: PCT/USO1/00670
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-30
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US-09-864-761-15294
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PRIOR FILING DATE: 2000-05-6
PRIOR PRILICATION NUMBER: US 09/632,366
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
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APPLICANT: Penn, Sha
APPLICANT: Rank, Da
APPLICANT: Hanzel,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local :
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR EPPLICATION NUMBER: US 60/180,312 PRIOR FILING DATE: 2000-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           73 TOCCTCCAGTTCCTGCGGCTCAGAGTCCCTGGCCCAGATCCCTCTG 119
69 TCCCTCCAGTTTCCTGCTGGGACTCGCCATTGCCCATATCCAGCNAG 23
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Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 4926
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(4803)
US-09-728-952-18
     ; Sequence 5, Application US/09893238
; Patent No. US20020150973A1
; GENERAL INFORMATION:
                                                                        RESULT 11
US-09-893-238-5/c
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US-09-728-952-18/c
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                   Similarity 60.6
59; Conservative
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APPLICANT: Zhou, Ping
APPLICANT: Liu, Chenghua
APPLICANT: Liu, Chenghua
APPLICANT: Asundi, Vinod
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Durrui
APPLICANT: Wang, Livrui
APPLICANT: Usundi, Manusha L.
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: NO. USZ0020111302A1el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 799
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TYPE: DNA

TYPE: DNA

ORGANISM: HOMO SAPIENS

FEATURE:

OTHER INFORMATION: MAP TO ACC24196.2

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.8

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.42

US-09-864-761-15294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/728,952
CURRENT FILING DATE: 2000-11-30
NUMBER OF SEQ ID NOS: 101
SOFTWARE: pt_FL_genes Version 2.0
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 15294
4220 TTGTAGCTCCCATCCGTGTTGGTGCAAGTGCCTCCCC 4184
                                                                                                         4280 TCGATGTCCTCACAGGCAGTGCCCTTGGCCCGTCAGCTCATGCCCAGGGGGACACTGGCAC 4221
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                                71 TT-TCGCTCCAGTTCCTGGGGCTCAGAGTCCCTGGCC 106
                                                                                                                                                            11 TTGCTCTCCTCAGAGGCCCCTGCGAGTGGAGTTCAGCTTTTCCTCATGGGGCAAATCTYAC 70
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                                                                                                                                                                                                                   Score 27; DB 10;
Pred. No. 11;
1; Mismatches 36
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND TITLE OF INVENTION: DIAGNOSIS OF BODY WEIGHT DISORDERS, INCLUDING OBESITY FILE REFERENCE: 7853-237

CURRENT APPLICATION NUMBER: U2/09/893,238

CURRENT FILING DATE: 2001-06-27

PRIOR APPLICATION WHERE: 09/245,041

PRIOR FILING DATE: 1999-02-05

PRIOR PRIOR DATE: 1999-02-05

PRIOR APPLICATION WUMBER: 60/093,630

PRIOR PILING DATE: 1998-07-21

PRIOR APPLICATION NUMBER: 60/104,978

PRIOR FILING DATE: 1998-07-21

PRIOR PRIOR FILING DATE: 1998-07-21

PRIOR DATE: 1998-07-21

PRIOR PRIO
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CURRENT APPLICATION NUMBER: US/09/789,561
CURRENT APPLICATION NUMBER: US/09/789,561
CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: PCT/US00/24008
PRIOR APPLICATION NUMBER: 60/152,317
PRIOR APPLICATION NUMBER: 60/152,317
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 60/152,315
PRIOR FILING DATE: 1999-09-03
PRIOR FILING DATE: 1999-09-03
NUMBER OF SEQ ID NOS: 194
SOFTMARE: PATENTIN VET. 2.0
SEQ ID NO 57
LENGTH: 1147
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-789-561-57
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                               RESULT 13
US-09-764-864-1714/c
; Sequence 1714, Application US/09764864
; Patent No. US20020132753A1
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; ORGANISM: Mus musculus
US-09-893-238-5
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Best Local S
Matches 48
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Patent No. US20020064818A1
GENERAL INFORMATION:
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TITLE OF INVENTION:
FILE REFERENCE: PZO4
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INFORMATION
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l Similarity 73.9%;
34; Conservative (
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; LOCATION: (2577)
; OTHER INFORMATION: n equals a,t,g, or
US-09-764-864-1714
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                                                                                                        PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR APPLICATION NUMBER: PCT/USO1/00661
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PRIOR FILING DATE: 2000-02-04
PRIOR PILICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 69/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRIOR
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PRIOR APPLICATION 2001-05-99
PRIOR APPLICATION OF THE CONTROL OF THE 
PRIOR APPLICATION NUMBER: PCT/US01/00670
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PRIOR FILING DATE: 2001-01-30
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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les 43; Conserv
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Chen, Wensheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              David R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22.1%; Score 26.8; D 59.7%; Pred. No. 19; tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 10; Length 15980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 5516
LENGTH: 472
TYPE: DNA
ORGANISM: Homo sapiens
APPLICANT: ELESTORI, FLUCIMENT APPLICANT: GENE LOSIG, INC.
TITTLE OF INVENTION: Molecular Toxicology Modeling
FILE REFERENCE: 44921-5038-US
CURRENT APPLICATION NUMBER: US/09/917,800A
CURRENT FILING DATE: 2001-07-31
PRIOR APPLICATION NUMBER: US 60/222,040
PRIOR FILING DATE: 2000-07-31
PRIOR APPLICATION NUMBER: US 60/222,880
PRIOR REILING DATE: 2000-01-02
PRIOR APPLICATION NUMBER: US 60/290,029
PRIOR APPLICATION NUMBER: US 60/290,645
PRIOR APPLICATION NUMBER: US 60/290,645
PRIOR APPLICATION NUMBER: US 60/292,336
PRIOR APPLICATION NUMBER: US 60/292,336
PRIOR FILING DATE: 2001-05-15
PRIOR PRILING DATE: 2001-05-22
PRIOR PRILING DATE: 2001-06-6-6
PRIOR PILING DATE: 2001-06-6-6
PRIOR APPLICATION NUMBER: US 60/292,798
PRIOR APPLICATION NUMBER: US 60/293,459
PRIOR APPLICATION NUMBER: US 60/298,884
PRIOR FILING DATE: 2001-06-19
PRIOR APPLICATION NUMBER: US 60/303,459
PRIOR APPLICATION NUMBER: US 60/295,798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:

OTHER INFORMATION: MAP TO ACO04613.1

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL - 2.2

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 1.7

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 1.8

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL - 1.8

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL - 2

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL - 2

OTHER INFORMATION: EXPRESSED IN BLART, SIGNAL - 2

OTHER INFORMATION: EXPRESSED IN BLART, SIGNAL - 1.7

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 1.7

OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 1.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
US-09-917-800A-61/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Mendrick, Donna
APPLICANT: Porter, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 61, Application US/09917800A Patent No. US20020119462A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Johnson, Kory
Castle, Arthur
Elashoff, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 7.2;
1; Mismatches 20;
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